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SEQUENCE LISTING

<110> diaDexus, Inc. Macina, Roberto Turner, Leah Sun, Yongming Rodriguez, Maria

<120> Compositions, Splice Variants and Methods Relating to Colon Specific Genes and Proteins

<130> DEX-0449

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<151> 2002-12-04

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| gaagtggage ccatcateet eteegtggge gaggaagatg aggteteeat caaggagge   | a 1200  |
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240

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| 420    | ccaggtcgcg | aaggcctgaa | ccccagggc  | actgcagtgt | ttcgccttgg | caagtctcct |
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| 1980   | cctgttgccc | gcccttccta | cttagacctg | cccagctcac | gggctgcacc | cctctctcta |
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1140

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| tttaagctgt | gtacagtgaa | aattgtcttt | actgtatttt | tgttctctgg | taatgtaata | 1440 |
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| agcatgatgg | tgccttctat | taatacatca | ttccagtctt | gctggtaatt | ttgtacagta | 1500 |
| tagtgtatga | attgctgtgc | tgcaaagcca | aacagctgca | aaatgttgaa | aaatcatcga | 1560 |
| aatgtataaa | aattgcagta | tctttaaaat | cagtaaaatg | gactagcata | ttatttatct | 1620 |
| tgttcttcag | ttaacaactt | tgtgttctct | gtgggaggga | gggagtcctg | tgtgtttgtg | 1680 |
| gggagaggga | aggaggaagt | cagttatttg | agtaagcctc | tagttgactt | ttctcttage | 1740 |
| ctgaatgtgg | acgttgaaac | atatcacttc | agggcttgga | aaagtcagtc | aacttgacgt | 1800 |
| acatttttag | tgacatttta | aaagcagtca | gattctataa | atggcaagta | agcctgaagt | 1860 |
| gaggatactg | caattttcgg | agaaaagaac | agcagetett | taagtgtttg | cattttctat | 1920 |
| ttggggggca | gggaactgtc | attcattttg | cacaattctt | gaactgatgt | cagcacccga | 1980 |
| gtggctcctg | aatttaagtc | tgggacgaca | tottttattt | ttacatgaat | ctttaaacaa | 2040 |
| ttctgtgagc | aaagtttgta | gctgctggat | tattgtctgt | ctttatagca | agttccagta | 2100 |
| aaccacaagt | atggcaaagc | ttatccaatt | ttatgcttgg | agcagtcagt | acataccagi | 2160 |
| ttctgatgtt | tcaggcagga | gtggggtaaa | taagtgtgac | cacttaaagc | tgctcgttag | 2220 |
| catggaagac | ttctccattc | tatotttgta | aaacagacaa | gatatgcact | tgacatagta | 2230 |
| gcaaattggt | tctgaattat | gcaactgttt | gctatttagt | aaactagcaa | atgatgcatg | 2340 |
| tattttgttt | ttcatgtact | gggcaatatg | agtaaaatct | gtcccttttt | ccccctttga | 2400 |
| atgaggtctt | ccatgtttga | gggaaagtct | tgcactattg | catatattt  | ggggacacag | 2460 |
| attttcatag | tttccatttt | tggggggctt | aaggattttt | tttttttctg | tttgaaacag | 2520 |
| ttttatactt | tctgatatag | tacttgaaat | tcttaccaga | aaattacttt | ggagttttga | 2580 |
| agcctttatt | aatactactt | ttaaagaagc | agttgtttta | ttgtcaatgt | ttttttccc  | 2640 |
| ccaagcatat | tttcttgtat | ttctgtttcc | atatatatat | atatatataa | atttccaatt | 2700 |
| caggatattg | ccctgccatc | catgaaaact | gttctggcac | caaaagtaat | gacaaatgtt | 2760 |
| aagtgtaata | atagaaaagt | agagcaaaga | gccattcagc | ttcagtcttt | acataccatg | 2820 |
| aataaaacat | taaaacatca | tatggagaag | tttacatggt | gattgttcac | ctgcagtact | 2880 |
| gtggagtttt | aacattttgt | cctcttttca | gtgaaacaga | gtaaaaatat | tcatctacca | 2940 |
| ttactgttat | ttgctgattt | tgttttattt | tttgatggta | atattctatc | cttatgacac | 3000 |
| tattgcaacc | aaattggctt | taccatcttg | gctttagtag | gtatagaaga | caatggatta | 3060 |
| ccatctttat | tgctgtaatg | tgttaagcat | tatatgctag | tagaatctag | tttaattgtt | 3120 |
| tcaggtggaa | agtattcttt | gagtttccat | attgaatgtg | tttggactaa | acaaacaata | 3180 |
|            |            |            |            |            |            |      |

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| aactactgat gtctgcagca   | 3200  |
|---|-------|
| <210> 22<br><211> 627<br><212> DNA<br><213> Homo sapien                       |       |
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| getttaacag gettegeetg tgetteetgt tteetettt accaaggace egecaacatg              | 120   |
| ggccgcgttc gcaccaaaac cgtgaagaag gcggcccggg tcatcataga aaagtactac             | 180   |
| acgcgcctgg gcaacgactt ccacacgaac aagcgcgtgt gcgaggagat cgccattatc             | 240   |
| cccagcaaaa agctccgcaa caagatagca ggttacgtca cgcatctgat gaagcgaatt             | 300   |
| cagagaggcc cagtaagagg tatctccatc aagctgcagg aggaggagag agaaaggaga             | 360   |
| gacaattatg ttcctgaggt ctcagccttg gatcaggaga ttattgaagt agatcctgac             | . 420 |
| actaaggaaa tgctgaagct titggasttc ggcagtctgt scaaccttca ggtcattcat             | 480   |
| cccaactgta ggctgagtga cctgaaggtt ggacagactg cagttgggat gaatttcaaa             | 540   |
| acgcctcggg gacctgtttg aattttttct gtagtgctgt attattttca ataaatctgg             | 600   |
| gacaacagca aaaaaaaaa aaaaaaa  | 627   |
| <210> 23<br><211> 3187<br><212> DNA<br><213> Homo sapien                      |       |
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| gtottoogot ocagottogo coacttooco ttgocagogg ggtgggogog gagaagacot             | 180   |
| gccggagcca tggaggacga agtggtccgc tttgccaaga agatggacaa gatggtgcag             | 240   |
| aagaagaagg tgagegegge gageggegeg ggeegggagg gaggeeggge egeggteegg             | 300   |
| cccgaggggg ccgtcccctc cctccttctc cctcccctcc                                   | 360   |
| gageegggee egeegegeee gegteeeegg ggagggtgte egggeeeggg gegtegggte             | 420   |
| ccgcggcaac ctgcgttagg acccgctcgc tcttgtattt ccagtccctg ctcgctggtg             | 480   |
| ccgtgtaata aacctaatcg atttagggat tagggcacgc tctccgttga ctttaaactt             | 540   |
| togtgotagt aaaacottgg catttagott aggaattgma gogtagaaag tootcagaac             | 600   |
| ttgaaagaat.ggagtcttcc gtgatccgtg gccaacatct agaaagagca tcaaatagag             | 660   |
| qcttqtqatc qotattaqac atccqqctoq taaaatgtaa aacttctgtg tcttattttt             | 720   |

| gaagagggct | gaacgtccgc | ctggttatct | caactgtgct | acatgtttgc | Étagcgatta | 780  |
|------------|------------|------------|------------|------------|------------|------|
| aaaacgcggc | tggagcattg | gatttgctaa | aggagcttaa | gaatattoot | atgaccctgg | 840  |
| aattactgca | gtccacaaga | atcggaatgt | cagttaatgc | tattcgcaag | cagagtacag | 900  |
| atgaggaagt | tacatctttg | gcaaagtctc | tcatcaaatc | ctggaaaaaa | ttattagatg | 960  |
| ggccatcaac | tgagaaagac | cttgacgaaa | agaagaaaga | acctgcaatt | acategeaga | 1020 |
| acagccctga | ggcaagagaa | gaaagtactt | ccagcggcaa | tgtaagcaac | agaaaggatg | 1080 |
| agacaaatgc | tcgagatact | tatgtttcat | cctttcctcg | ggcaccaagc | acttctgatt | 1140 |
| ctgtgcggtt | gaagtgtagg | gagatgettg | ctgcagctct | tcgaacaggg | gatgactaca | 1200 |
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| aagatgcaaa | aaatccaaat | ttaaggaaaa | atgtcctctg | tgggaatatt | cctcctgact | 1380 |
| tatttgctag | aatgacagca | gaggaaatgg | ctagtgatga | gctgaaagag | atgcggaaaa | 1440 |
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| agttctgttg | agttggaaga | attggcaaaa | tatctggacc | attaagaaaa | cggattttgt | 1580 |
| aactagcttt | aaactaggcc | aagcaactag | ttttcctgca | aatcaaattt | ttaaagcaac | 1740 |
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| atcagtaggg | agaccatata | ataattttat | ggtacctgtt | tcaaaacata | ttttttctgt | 1860 |
| ttttataagt | aagttgatat | taattaaact | cttggcaata | tttcttcttt | cttaaaggaa | 1920 |
| aatatacctt | aactttttt  | cttttacact | gtgaaacata | cacagtagaa | attctgttac | 1980 |
| tctctgttat | taatacataa | atgaaaatac | attttttcc  | atattggcat | gtagctacaa | 2040 |
| atattaaagg | aggagaaaag | gtaatataat | tttaggttta | ccaaatatgg | tgtgtattca | 2100 |
| aataatactt | gaccagctta | tctaaaatgt | acataatttt | gaggtagctt | atgaatttga | 2160 |
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| cgtgatcatc | atttcttgta | atttcttgta | catgtatatt | acttgttctt | aatagatttt | 2280 |
| tggaaacaag | actttattga | gatcagtttg | gttttcctgt | taatttacct | gtrtgacttt | 234  |
| ataatgtgtt | ttagttttgc | agaagaacac | tgttgtagtt | tagaaggctt | ttcataaatc | 240  |
| ccctcatagg | caaagatgaa | aacttcccac | tattttttc  | ccctcttagg | aagacatact | 246  |
| CC322C222  | tatttaacat | cttactctac | tatacctate | gtaaacagtt | catgactaga | 252  |

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| ttttgattcg                                      | gaaatctata   | ctgaccaagg   | attaatctta   | aggattgtat   | aattcattaa   | 2580                     |
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| tgactttttc                                      | tgtcatgagt   | gagattgttg   | aacaaactga   | atatatgggc   | tatagcaagt   | 2700                     |
| agctttacag                                      | tacagatctt   | acaattaagt   | tttgcttttg   | ttaaagtgtg   | taccattttt   | 2760                     |
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| gcatacttta                                      | aaggccactg   | ttgcaaagtc   | tacattttat   | gctgaatctg   | cattctgtca   | 2880                     |
| ggcacccgta                                      | gaaagacctc   | agtacatgct   | ttgcactctc   | ctttgctccc   | tttttccaat   | 2940                     |
| ttcttattgc                                      | atatcatttt   | gttgtaatac   | agaaagcagc   | atttttaaat   | gtccgtgtta   | 3000                     |
| agaattggcc                                      | cactggtacc   | aactcacctc   | tattttgtca   | gttcatagtt   | gaagattttg   | 3060                     |
| ttttatttca                                      | aaaacaaagt   | acatttttga   | aataatgttt   | cagaataaaa   | taatctcact   | 3120                     |
| tttaagtgat                                      | ccattttaaa   | atttgtaatt   | caataaagtt   | ttttttgttg   | ttaaacataa   | 3180                     |
| aaaaaaa   |  |  |  |  |  | 3187                     |
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| cccgggagct                                      | cgggacgcag   | gcggggcttg   | tgctccgcgg   | gggcagggcg   | tagggtgggc   | 120                      |
| ctcctacctc                                      | ccctgatctc   | gcggtttgtt   | ccgtttcatt   | ggagcttccc   | ggaccgtgtg   | 180                      |
| ctcgacggtg                                      | ccctaggtgc   | cgtggggcca   | cacgcgagtc   | tgataagcac   | cctccccgg  | 240                      |
| aatcatgcgg                                      | tgctgtgagg   | cctagcgaag   | atgaagatag   | aatgcaaggt   | agaaagtgct   | 300                      |
| ggataccttt                                      | agaaagctgc   | aggactggtg   | cgatgggagt   | tgagacgtaa   | gaacctgccc   | 360                      |
| gtccgtaggg                                      | ctctggatgc   | tgctgaggcc   | cgaggcccct   | atggcagatt   | tgaaaattca   | 420                      |
| cccttgtaga                                      | gtcattcctg   | cctttgagcg   |  |  |  |                          |
| ttcggtggag                                      | 2 2  | ccccgageg  | gactcccttt   | taagagatct   | caagagagcg   | 480                      |
| ggagccggcg                                      | gccctgggtc   |  |  |  |  | 480<br>540               |
|   |  | tgcacagete   | acctccctgg   | gaactgctcg   | cccgagcgtc   |                          |
| ggacacaggg                                      | gccctgggtc   | tgcacagctc<br>gcagccggaa                               | acctccctgg<br>ggttgcagcc                               | gaactgctcg   | cccgagcgtc   | 540                      |
|   | gccctgggtc   | tgcacagctc<br>gcagccggaa<br>tgcagaatcc                 | acctccctgg<br>ggttgcagcc<br>acaggtcttt                 | gaactgctcg<br>gcaggagccc<br>cttgaggaaa                             | cccgagcgtc<br>cggaggccca<br>tctgtagaca                             | 540<br>600               |
| gaactttgtg                                      | gccctgggtc<br>ctggccccct<br>ctcttgctct               | tgcacagctc gcagccggaa tgcagaatcc atctagggaa            | acctccctgg ggttgcagcc acaggtcttt ggaacagaag            | gaactgctcg<br>gcaggagccc<br>cttgaggaaa<br>agtgtcgtct               | cccgagcgtc<br>cggaggccca<br>tctgtagaca<br>cctagaaatc               | 540<br>600<br>660        |
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900

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| agtgatccac ccaccttg  | gc cttccaaagt  | ggtgggatta  | tgggcaggag | cctccgtgcc | 960  |
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| gccagetect taccacag  | ag acgccgcgtg  | gaactcacta  | ctggcgatcg | cggacgcccc | 240  |
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| ggtgccaggc tcacccct  | ca gatacagcct  | cgtgggctgg  | ctcagtggcg | gccctgactc | 360  |
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| aaacctgcct ggctgtc   | cc ttccgggaca  | cggacagggg  | gtttcctggc | agttcctggt | 540  |
| cctcccttga gggctaag  | ggg tececeegtt | ctgtggccgc  | ctccagcaca | tccacccaga | 600  |
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300

| cgtgctgggc | cctgggcgga | ctccccggct | ggaggaccag | caggetetge | cctacacaag | 1200 |
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|                                    |              |            | cgggccctca |            |              | 540  |
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|   | ggcacgaaag  |            |            |            |            |            |
|   | cagtttcaag  |            |            |            |            | 360<br>420 |
|   | ccccggcgag  |            |            |            |            |            |
|   | gaatgacaag  |            |            |            |            | 480<br>540 |
|   |             |            | Jecugaat   |            | Jagaacetta | 240        |

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| tgctgcgctc | agccatgcgc | ctgtgcctgc | cgagtttcga | caagctggag | ttgctggagt | 720  |
| gcatccgccg | gctcatcgaa | gtggacaagg | actgggtccc | cgatgccgcc | ggcaccagcc | 780  |
| tctatgtgcg | gcctgtgctc | attgggaacg | agccctcgct | gggtgtcagc | cagcccacgc | 340  |
| gegegeteet | gttcgtcatt | ctctgcccag | tgggtgccta | cttccctgga | ggctccgtga | 900  |
| ccccggtctc | cctcctggcc | gacccagcct | tcatccgggc | ctgggtgggc | ggggtcggca | 960  |
| actacaagtt | aggtgggaat | tatgggccca | ccgtgttagt | gcaacaggag | gcactcaagc | 1020 |
| ggggctgtga | acaggtcctc | tggctgtatg | ggcccgacca | ccagctcacc | gaggtgggaa | 1080 |
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| cgctgaatgg | tgttatcctg | cctggagtgg | tcagacagag | tctactggac | atggctcaga | 1200 |
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| agtetttgge | tegggeaceg | cttgccaggt | ctgcccagtg | caccgaatcc | tgtacaaaga | 1560 |
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| ttggccccca | gcccctcgtc | gcgggttcag | gtccgcccat | tactccttgt | cgtgcggtca | 1920 |
| aggatacacc | ttggccccga | ttccggatct | ctccgttctc | aggccagacc | catggtgctg | 1980 |
| ccattaattt | tttttttt   | atetttaeta | caattttgaa | ataaaatgcc | aaagaacaca | 2040 |

<sup>&</sup>lt;210> 43

<sup>&</sup>lt;211> 1191

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 43

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tgtaacggtc cctgcgtgca tttatgcgct ctttggaccc tgtgaacaca ccgggatact 120

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77

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| ctgcttcata                               | tgtcccaaaa    | gagaaaaaaa | taaaggggac | aatgccaaca | tgctcaacaa | 300  |
| taaaggcttc                               | tttttcttat    | ttttttaata | caaaatacaa | gcaaaggata | cacatactta | 360  |
| aaacagagct                               | caggagcaga    | cacgcagtcc | tggaaaccct | tcaataaaag | caaagcagga | 420  |
| gtttgttttt                               | tctttgtcta    | tgcagataca | tacagagact | gggatatgta | aaaattaagt | 480  |
| atcacaaaag                               | accatcacac    | gattctacca | atgcatgttg | catctgtaat | tcacgaacat | 540  |
| ggtcaacaaa                               | atcatgttca    | cttcaacccc | atttcattta | aattaaagaa | aaaaaccttt | 600  |
| taaataaagt                               | ggttacattc    | aaactttaac | ttccttagta | ccatgctgca | gatttcagca | 660  |
| ctgttaaggt                               | attgcaagaa    | tgcccaaccc | tctggtgtct | gatcatgtat | ctagcaacat | 720  |
| tgcagtatga                               | agaaaagaga    | tgccccggtc | tcagcccatg | gactagttaa | tacagtgaag | 780  |
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| cttatttaca                               | aatcttgtaa    | caaatacttc | tggaggaaaa | agagaaaaga | attcactaag | 1080 |
| ttccagaaga                               | caagcttaat    | tgcagacgta | tacaaacaca | catcacacgt | aacaccacac | 1140 |
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| ggccgcggcc                               | tccgcgggcc    | caggggcctc | caggccccgt | gaccacegga | ccgagcaccc | 180  |
| tggaacggcc                               | ccagcttggc    | ctcgggacag | tccgcgcgct | caccgactcc | ctggtgaacg | 240  |
| ccgcttggcc                               | gccgccgccg    | ccgcaggacc | cacgagaagc | ggaaaccggc | gcaaggaccc | 300  |
| ggtccccccg                               | gcgccgtacg    | tggagtgagc | ctgcggctcc | gccccgcgcg | ctccgattgg | 360  |

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PCT/US2003/040131

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540 600

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| tccctctgtt | cagtggattc | ttgactacat | ataggtcata | tatttcaaaa | aataatgcct | 960  |
| agctatttct | actttgaaat | catgactaaa | gccaaaccac | aaccacagca | aagataacct | 1020 |
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| acttgtaatc | ccagcacttt | gagaggccga | gatgggcaga | tcacctgagg | tcaggggttc | 1140 |
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|   |               |            |            | gtaaaaccag |            | 130  |
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| <pre>&lt;211&gt; 2435 &lt;212&gt; DNA &lt;213&gt; Homo sapien  &lt;400&gt; 57 ccccttccag tccctgctcc ttcaatagac agaagcacga aacctcccct agatcgttca ttagctccgt ttgatagaga acccttcaca ctaggggagg ttgcgtgctt ctgtctattg aaggagcacg aaacctcccc tagatcgttc attagctccg tttgatagag aacccttcac actaggaaag aaaccaccat tttctgacaa gccctcgatt ccagcgggaa ggtcactcgg ggagcattta cccaagattc aaaagcctcc tttaccaccg accacggaaa gacatgaaag gagcagccc ctgccaggga agaagccacc tgtgccaaag catggatggg gaccagacag aagagagaat gatgaagatg atgtgcatca gagacctttg ccccagccag cactacttcc tatgagctcc aacactttcc cttcaagatc tactaagcca agtcccatga accctcccc atcctccac atgcctggag cattctcaga aagtaacagc agttttccac agagtgcctc</pre>  | 120<br>180<br>240<br>300<br>360<br>420<br>480<br>540        |
| <pre>&lt;211&gt; 2435 &lt;212&gt; DNA &lt;213&gt; Homo sapien  &lt;400&gt; 57 ccccttccag tccctgctcc ttcaatagac agaagcacga aacctcccct agatcgttca ttagctccgt ttgatagaga acccttcaca ctaggggagg ttgcgtgctt ctgtctattg aaggagcacg aaacctcccc tagatcgttc attagctccg tttgatagag aacccttcac actaggaaag aaaccaccat tttctgacaa gccctcgatt ccagcgggaa ggtcactcgg ggagcattta cccaagattc aaaagcctcc tttaccaccg accacggaaa gacatgaaag gagcagccc ctgccaggga agaagccacc tgtgccaaag catggatggg gaccagacag aagagagaat gatgaagatg atgtgcatca gagacctttg ccccagccag cactacttcc tatgagctcc aacactttcc cttcaagatc tactaagcca agttccatga accctcccc atcctctcac atgcctggag cattetcaga aagtaacagc agttttccac agagtgcctc cctgccacca tacttctctc aaggccctag caacagacca cctatcagag ccgaaggcag</pre>   | 120<br>180<br>240<br>300<br>360<br>420<br>480<br>540        |
| <pre>&lt;211&gt; 2435 &lt;212&gt; DNA &lt;213&gt; Homo sapien </pre> <pre>&lt;400&gt; 57 ccccttccag tccctgctcc ttcaatagac agaagcacga aacctcccct agatcgttca ttagctccgt ttgatagaga acccttcaca ctaggggagg ttgcgtgctt ctgtctattg aaggagcacg aaacctcccc tagatcgttc attagctccg tttgatagag aacccttcac actaggaaag aaaccacat tttctgacaa gccctcgatt ccagcgggaa ggtcactcgg ggagcattta cccaagattc aaaagcctcc tttaccaccg accacggaaa gacatgaaag gagcagccc ctgccagga agaagccacc tgtgccaaag catggatgg gaccagacag aagagagaat gatgaagatg atgtgcatca gagacctttg ccccagccag cactacttcc tatgagctcc aacactttcc cttcaagatc tactaagcca agtcccatga accctcccc atcctctcac atgcctggag cattctcaga aagtaacagc agttttccac agagtgcctc cctgccacca tacttctctc aaggccctag caacagacca cctatcagag ccgaaggcag aaacttcccc ttgccacttc caaacaaacc tcggcccca tccccgcgg aggaagagaa</pre> | 120<br>180<br>240<br>300<br>360<br>420<br>480<br>540<br>600 |

| acttgtaaat | catttctgaa   | atttataagg  | tttccagctg   | acaattttct  | gccaagttaa   | 900  |
|------------|--------------|-------------|--------------|-------------|--------------|------|
| catttcaaaa | atgaaaaaca   | aaatcaatgt  | gtattttaga   | gtatcatcta  | gatctgcgcc   | 960  |
| cctttgtaaa | tggattctat   | ggtggttaag  | tcatctagaa   | acaccctggg  | agcaaggttc   | 1020 |
| catcctacta | cttcctaatt   | acatctgatc  | tgctttattt   | ttgtgtcctc  | tttatactag   | 1080 |
| ctgtttctaa | ctggggggcg   | gttatggaat  | gtttcttagt   | gacagctggg  | gcctgccaat   | 1140 |
| ctcattcatt | gttttctact   | tctgcacgtg  | catcagagac   | ctttgcccca  | gccagcacta   | 1200 |
| cttcctatga | gctccaacac   | tttcccttca  | agatctacta   | agccaagtcc  | catgaaccct   | 1260 |
| ctcccatcct | ctcacatgcc   | tggagcattc  | tcagaaagta   | acagcagttt  | tccacagagt   | 1320 |
| gcctccctgc | caccatactt   | ctctcaaggc  | cctagcaaca   | gaccacctat  | cagagccgaa   | 1380 |
| ggcagaaact | tccccttgcc   | acttccaaac  | aaacctegge   | ccccatcccc  | cgcggaggaa   | 1440 |
| gagaattcat | taaatgaaga   | gtggtacgtt  | tcttatatta   | cccgaccaga  | ggcagaagct   | 1500 |
| gctcttagaa | agataaacca   | ggatggcaca  | tttctggtca   | gagacagete  | taaaaaaaca   | 1560 |
| acaaccaatc | catatgtcct   | catggtgttg  | tacaaagata   | aagtttacaa  | catccagatc   | 1620 |
| cgttatcaga | aggaaagtca   | agtttacttg  | ttgggaactg   | gactccgagg  | gaaagaggac   | 1680 |
| tttctgtctg | tgtcagatat   | tattgactac  | ttcaggaaaa   | tgccacttct  | gctcattgat   | 1740 |
| gggaaaaacc | gaggttccag   | ataccagtgo  | acattaacgc   | atgctgcagg  | gtacccatag   | 1800 |
| caagttatag | ; ccgagcaaat | gaaccgtcct  | cctgcctctg   | ttgccaacac  | gagatcaatc   | 1860 |
| agccttggtc | aatggacaaa   | cacttaggad  | : tgaactgaac | ccctccccat  | gaacacaagg   | 1920 |
| gttttatcct | ttcctttaaa   | aacagtgttt  | gaaatgaaga   | ctgtcaacta  | teccataatt   | 1980 |
| tatttattct | : tcttcaatgt | tigtaaagtg  | g catgagtcat | gttcacactt  | gaagtctagt   | 2040 |
| agtgcactgt | aataattcat   | ttttaaaag   | g attatttaat | gcccatttca  | a aaatacagta | 2100 |
| gtttacacag | g ctacagaaa  | aatttgggg   | aagttttaaa   | acactgaaad  | agtaatagtt   | 2160 |
| attggtgtca | a cataaaact  | g atttgtttt | tacagccaaa   | cctctgtcag  | g tcagaggcat | 2220 |
| tcattagtt  | t tatacatgta | a atttgaaaa | cactaaacct   | cgttttctca  | a gcagcaataa | 2280 |
| tttaagagg  | c ttcaaaaata | a taatttcac | t cttatttagt | atttttcc    | gggggcattt   | 2340 |
| ttacgtaat  | t tttttatga  | a aagacaaat | g catgttgaga | a taacttctg | g gattaaaata | 2400 |
| gtcttttgc  | t ttaaaaaac  | c tegeegega | c cacgc      |             |              | 2435 |

<sup>&</sup>lt;210> 58 <211> 3612 <212> DNA <213> Homo sapien

<sup>&</sup>lt;400> 58

|            |              |              | 100        |              |              |          |
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| cgcgcccatc | gggtaccgag   | gcacgcgccg   | ggcgtcacgt | gcgtttcgcg   | gcgagcggaa   | 120      |
| atgacgcgag | ttgtgtgagc   | cgccagtatg   | gccgggctat | ggcggcgagc   | actggctacg   | 180      |
| tgcgactgtg | gggagcggcg   | cggtgctggg   | tgctgcggcg | gccgatgctg   | geegeegeeg   | 240      |
| gggggcgggt | tcccactgca   | gcaggagcgt   | ggttgctccg | aggccagcgg   | acctgcgacg   | 300      |
| cctctcctcc | ttgggcactg   | tggggccgag   | gcccggcaat | tgggggccaa   | tggcgggggt   | 360      |
| tttgggaagc | gagcagccgc   | ggcggaggcg   | cattctcggg | gggcgaggac   | gcctccgagg   | 420      |
| gcggcgcgga | ggaaggagcc   | ggcggcgcgg   | ggggcagcgc | gggcgccggg   | gaaggcccgg   | 480      |
| tcataacggc | gctcacgccc   | atgacgatcc   | ccgatgtgtt | teegeacetg   | ccgctcatcg   | 540      |
| ccatcacccg | caacccggtg   | ttcccgcgct   | ttatcaagat | tatcgaggtt   | aaaaataaga   | 600      |
| agttggttga | gctgctgaga   | aggaaagttc   | gtctcgccca | gccttatgtc   | ggcgtattta   | 660      |
| taaagagaga | tgacagcaat   | gagtcggatg   | tggtcgagag | cctggatgaa   | atctaccaca   | 720      |
| cggggacgtt | tgcccagatc   | catgagatge   | aggaccttgg | ggacaagctg   | cgcatgatcg   | 780<br>· |
| tcatgggaca | cagaagagte   | catatcagca   | gacagctgga | ggtggagccc   | gaggagccgg   | 840      |
| aggcggagaa | caagcacaag   | ccccgcagga   | agtcaaagcg | gggcaagaag   | gaggcggagg   | 900      |
| acgagctgag | cgccaggcac   | ceggeggage   | tggcgatgga | gcccacccct   | gageteeegg   | 960      |
| ctgaggtgct | catggtggag   | gtagagaacg   | ttgtccacga | ggacttccag   | gtcacggagg   | 1020     |
| aggtgaaagc | cctgactgca   | gagatcgtga   | agaccatccg | ggacatcatt   | gccttgaacc   | 1080     |
| ctctctacag | ggagtcagtg   | ctgcagatga   | tgcaggctgg | ccagcgggtg   | gtggacaacc   | 1140     |
| ccatctacct | gagcgacatg   | ggcgccgcgc   | tcaccggggc | cgagtcccat   | gagctgcagg   | 1200     |
| acgtcctgga | agagaccaat   | attcctaagc   | ggctgtacaa | ggecetetee   | ctgctgaaga   | 1260     |
| aggaatttga | actgagcaag   | ctgcagcagc   | gcctggggcg | ggaggtggag   | gagaagatca   | 1320     |
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| tggacaacca | ctcctcggag   | ttcaatgtca   | cccgcaacta | cctagactgg   | ctcacgtcca   | 1560     |
| tecettgggg | r caagtacago | aacgagaacc   | tggacctggc | gcgggcacag   | gcagtgctgg   | 1620     |
| aggaagacca | ı ctacggcatg | gaggacgtca   | agaaacgcat | cctggagttc   | attgccgtta   | 1680     |
| gccagctccg | g cggctccacc | : cagggcaaga | tectetgett | ctatggcccc   | cctggcgtgg   | 1740     |
| gtaagaccag | g cattgctcgc | : tecategeee | gcgccctgaa | a ccgagagtac | : ttccgcttca | 1300     |
| gcgtcgggg  | g catgactgac | : gtggctgaga | tcaagggcca | a caggeggaee | : tacgtgggcg | 1860     |
|            |              |              |            |              |              |          |

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| tcatcgacga | ggtgcgtgcg   | gagaggcccg   | gagaccccct   | aacccggcaa   | tgccgtgtgc | 1980 |
| aagatgaagc | cgagccttcc   | gggtcctagg   | atccccgagc   | caactcaggg   | gccatcgccg | 2040 |
| tgatgctggc | cctgatggca   | cggacgggtc   | cctcccaccc   | acagccacgc   | ccagcaggct | 2100 |
| gcccgtccaa | ggcagggtgt   | gggtgccgct   | ctcactcggg   | gtctccccag   | ccctgaccct | 2160 |
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| gacttgtcca | aggtgggggg   | cctgtctggg   | gcctgggctt   | gctggggagg   | gtaagcggcc | 2400 |
| agtcccccag | caccttgccg   | ccgcccccag   | gtgctgttca   | tctgcacggc   | caacgtcacg | 2460 |
| gacaccatcc | ccgagccgct   | gcgagaccgt   | atggagatga   | tcaacgtgtc   | gggctacgtg | 2520 |
| gcccaggaga | agctggccat   | tgcggagcgc   | tacctggtgc   | cccaggctcg   | cgccctgtgt | 2580 |
| ggcttggatg | agagcaaggc   | caagetgtca   | tcggacgtgc   | tgacgctgct   | catcaagcag | 2640 |
| tactgccgcg | agagcggtgt   | ccgcaacctg   | cagaagcaag   | tggagaaggt   | gttacggaaa | 2700 |
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| acaggccago | : tgggggaggt | gatgaaggag   | agcgcccgca   | tagcctacac   | cttcgccaga | 3000 |
| gccttcctca | tgcagcacgo   | ccccgccaat   | gactacctgg   | tgacetcaca   | catccacctg | 3060 |
| catgtgcccg | agggcgccac   | ccccaaggac   | ggcccaagcg   | caggetgeac   | catcgtcacg | 3120 |
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| aagcgcgcag | g gggtgacgtg | ; categteetg | r ccagccgaga | acaagaagga   | cttctacgac | 3300 |
| ctggcagcct | tcatcaccga   | gggcctggag   | gtgcacttcg   | , tggaacacta | ccgggagatc | 3360 |
| ttcgacatc  | g cettecegga | a cgagcaggca | . gaggegetgg | g ccgtggaacg | gtgacggcca | 3420 |
| ccccgggact | t acsaacaac  | g gatgtcaggo | cctgtctggg   | g ccagaactga | gcgctgtggg | 3480 |
| gagcgcgcc  | c ggacctggca | a giggagecad | cgagcgagca   | a geteggteea | gtgacccaga | 3540 |
| tcccaggga  | c ctcagtcgg  | c ttaatcagag | g tgtggcatag | g aagctattta | atgattaaag | 3600 |
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1560

1620

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| gctacctggt                             | gccccaggct | cgcgccctgt | gtggcttgga | tgagagcaag | gccaagctgt | 1740 |
| catcggacgt                             | gctgacgctg | ctcatcaagc | agtactgccg | cgagagcggt | gtccgcaacc | 1800 |
| tgcagaagca                             | agtggagaag | gtgttacgga | aatcggccta | caagattgtc | agcggcgagg | 1860 |
| ccgagtccgt                             | ggaggtgacg | cccgagaacc | tgcaggactt | cgtggggaag | cccgtgttca | 1920 |
| ccgtggagcg                             | catgtatgac | gtgacaccgc | ccggcgtggt | catggggctg | gcctggaccg | 1980 |
| caatgggagg                             | ctccacgctg | tttgtggaga | catccctgag | acggccacag | gacaaggatg | 2040 |
| ccaagggtga                             | caaggatggc | agcctggagg | tgacaggeca | gctgggggag | gtgatgaagg | 2100 |
| agagcgcccg                             | catageetae | accttcgcca | gagccttcct | catgcagcac | gcccccgcca | 2160 |
| atgactacct                             | ggtgacctca | cacatccacc | tgcatgtgcc | cgagggcgcc | acccccaagg | 2220 |
| acggcccaag                             | cgcaggctgc | accatcgtca | cggccctgct | gtccctggcc | atgggcaggc | 2280 |
| ctgtccggca                             | gaatctggcc | atgactggcg | aagtctccct | cacgggcaag | atcctgcctg | 2340 |
| ttggtggcat                             | caaggagaag | accattgcgg | ccaagcgcgc | aggggtgacg | tgcatcgtcc | 2400 |
| tgccagccga                             | gaacaagaag | gacttctacg | acctggcagc | cttcatcacc | gagggcctgg | 2460 |
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| cagaggcgct                             | ggccgtggaa | cggtgacggc | caccccggga | ctgcaggcgg | cggatgtcag | 2580 |
| gccctgtctg                             | ggccagaact | gagcgctgtg | gggagcgcgc | ceggaeetgg | cagtggagcc | 2640 |
| accgagcgag                             | cageteggte | cagtgaccca | gatcccaggg | acctcagtcg | gcttaatcag | 2700 |
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| cgcgcccatc | gggtaccgag | gcacgcgccg | ggcgtcacgt | gcgtttcgcg | gcgagcggaa | 120 |
| atgacgcgag | ttgtgtgagc | cgccagtatg | gccgggctat | ggcggcgagc | actggctacg | 180 |
| tgcgactgtg | gggagcggcg | cggtgctggg | tgctgcggcg | gccgatgctg | gccgccgccg | 240 |
| gggggcgggt | tcccactgca | gcaggagcgt | ggttgctccg | aggccagcgg | acctgcgacg | 300 |
| cetetectee | ttgggcactg | tggggccgag | gcccggcaat | tgggggccaa | tggcgggggt | 360 |
| tttgggaagc | gagcagccgc | ggcggaggcg | cattctcggg | gggcgaggac | gcctccgagg | 420 |
| gcggcgcgga | ggaaggagcc | ggcggcgcgg | ggggcagcgc | gggcgccggg | gaaggcccgg | 480 |

| tcataacggc | gctcacgccc | atgacgatcc | ccgatgtgtt | tccgcacctg | ccgctcatcg | 540  |
|------------|------------|------------|------------|------------|------------|------|
| ccatcacccg | caacccggtg | ttcccgcgct | ttatcaagat | tatcgaggtt | aaaaataaga | 600  |
| agttggttga | gctgctgaga | aggaaagttc | gtctcgccca | gccttatgtc | ggcgtctttc | 660  |
| taaagagaga | tgacagcaat | gagtcggatg | tggtcgagag | cctggatgaa | atctaccaca | 720  |
| cggggacgtt | tgcccagatc | catgagatgc | aggaccttgg | ggacaagctg | cgcatgatcg | 780  |
| tcatgggaca | cagaagagtc | catatcagca | gacagctgga | ggtggagccc | gaggagccgg | 840  |
| aggcggagaa | caagcacaag | ccccgcagga | agtcaaagcg | gggcaagaag | gaggcggagg | 900  |
| acgagctgag | cgccaggcac | ccggcggagc | tggcgatgga | gcccacccct | gageteeegg | 960  |
| ctgaggtgct | catggtggag | gtagagaacg | ttgtccacga | ggacttccag | gtcacggagg | 1020 |
| aggtgaaagc | cctgactgca | gagatcgtga | agaccatccg | ggacatcatt | gccttgaacc | 1080 |
| ctctctacag | ggagtcagtg | ctgcagatga | tgcaggctgg | ccagcgggtg | gtggacaacc | 1140 |
| ccatctacct | gagegaeatg | ggcgccgcgc | tcaccggggc | cgagtcccat | gagctgcagg | 1200 |
| acgtcctgga | agagaccaat | attcctaagc | ggctgtacaa | ggccctctcc | ctgctgaaga | 1260 |
| aggaatttga | actgagcaag | ctgcagcagc | gcctggggcg | ggaggtggag | gagaagatca | 1320 |
| agcagaccca | ccgtaagtac | ctgctgcagg | agcagctaaa | gatcatcaag | aaggagctgg | 1380 |
| gcctggagaa | ggacgacaag | gatgccatcg | aggagaagtt | ccgggagcgc | ctgaaggagc | 1440 |
| tegtggteee | caagcacgtc | atggatgttg | tggacgagga | gctgagcaag | ctgggcctgc | 1500 |
| tggacaacca | ctcctcggag | ttcaatgtca | cccgcaacta | cctagactgg | ctcacgtcca | 1560 |
| tcccttgggg | caagtacagc | aacgagaacc | tggacctggc | gcgggcacag | gcagtgctgg | 1620 |
| aggaagacca | ctacggcatg | gaggacgtca | agaaacgcat | cctggagttc | attgccgtta | 1680 |
| gccagctccg | cggctccacc | cagggcaaga | tcctctgctt | ctatggcccc | cctggcgtgg | 1740 |
| gtaagaccag | cattgctcgc | tccatcgccc | gcgccctgaa | ccgagagtac | ttccgcttca | 1300 |
| gcgtcggggg | catgactgac | gtggctgaga | tcaagggcca | caggcggacc | tacgtgggcg | 1860 |
| ccatgcccgg | gaagatcatc | cagtgtttga | agaagaccaa | gacggagaac | cccctgatcc | 1920 |
| tcatcgacga | ggtggacaag | atcggccgag | gctaccaggg | ggacccgtcg | tcggcactgc | 1980 |
| tggagctgct | ggacccagag | cagaatgcca | acttcctgga | ccactacctg | gacgtgcccg | 2040 |
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| ccattgcgga | gcgctacctg | gtgccccagg | ctcgcgccct | gtgtggcttg | gatgagagca | 2220 |
| aggccaagct | gtcatcggac | gtgctgacgc | tgctcatcaa | gcagtactgc | cgcgagagcg | 2280 |
| gtgtccgcaa | cctgcagaag | caagtggaga | aggtgttacg | gaaatcggcc | tacaagattg | 2340 |

| tcagcggcga                                      | ggccgagtcc   | gtggaggtga    | cgcccgagaa | ecegeaggae   | cccgcgggga | 2400 |
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| agcccgtgtt                                      | caccgtggag   | cgcatgtatg    | acgtgacacc | gcccggcgtg   | gtcatggggc | 2460 |
| tggcctggac                                      | cgcaatgggt   | gagcgtggtg    | gcgggagaag | accccaaagc   | cattcccatt | 2520 |
| tctacccgag                                      | aacaagcagg   | tcccaccttg    | tacacctgtg | cagtgggcct   | caggtggctc | 2580 |
| tgaacggcct                                      | ctggcggggg   | gtggtgtgct    | gggagcccag | ggggtctggg   | ctgagggagg | 2640 |
| ggaaggctgt                                      | gacacatggg   | ggtggcagca    | caccatgggc | cttatggatt   | tgggtgccac | 2700 |
| tttgataggc                                      | ttacaggtaa   | ggacaagctc    | atccagatgg | gc           |            | 2742 |
| <210> 61<br><211> 211<br><212> DNA<br><213> Hom |              |               |            |              |            |      |
| <400> 61<br>tgttgtactg                          | gctggagtac   | agtggtgcag    | tctcagctca | ctgcaacctc   | cgcctctcgg | 60   |
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| acgcccagct                                      | aattttgtat   | ttttagtaaa    | gatggtgttt | ctccatgttg   | gtcaggctgg | 180  |
| tctcgaactc                                      | ccgacctcag   | gtgatccgcc    | cacctcggcc | tcccaaagtg   | ctggcatgac | 240  |
| agagctagag                                      | ctcgggccca   | gccccaggct    | gcagcccatt | cgcaggcacc   | cgaaagaact | 300  |
| tccccagtat                                      | ggtggtcctg   | gaaaggacat    | ttttgaagat | caactatatc   | ttcctgtgca | 360  |
| ttccgatgga                                      | . atttcagttc | atcagatgtt    | caccatggcc | accgcagaac   | accgaagtaa | 420  |
| ttccagcata                                      | gcggggaaga   | tgttgaccaa    | ggtggagaag | aatcacgaaa   | aggagaagtc | 480  |
| acagcaccta                                      | gaaggcagcg   | cctcctcttc    | actctcctct | gattagatga   | aactgttacc | 540  |
| ttaccctaaa                                      | cacagtattt   | ctttttaact    | tttttatttg | taaactaata   | aaggtaatca | 600  |
| cagccaccaa                                      | cattccaago   | taccctgggt    | acctttgtgc | agtagaagct   | agtgagcatg | 660  |
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| aagaaaggct                                      | ggggatattt   | gggttggctt    | ggttttgatt | ttttgcttgt   | ttgtttgttt | 780  |
| tgtactaaaa                                      | a cagtattato | ttttgaatat    | cgtagggaca | taagtatata   | catgttatcc | 840  |
| aatcaagat                                       | g gctagaatgg | tgcctttctg    | agtgtctaaa | acttgacacc   | cctggtaaat | 900  |
| ctttcaacac                                      | acttccactg   | cctgcgtaat    | gaagttttga | ttcattttta   | accactggaa | 960  |
| tttttcaatq                                      | g ccgtcatttt | cagttagatg    | attttgcact | ttgagattaa   | aatgccatgt | 1020 |
| ctatttgatt                                      | agtcttattt   | tttatttt      | acaggcttat | : cagtctcact | gttggctgtc | 1080 |
| attgtgaca                                       | a agtcaaataa | acccccaagg    | acgacacaca | gtatggatca   | catattgttt | 1140 |
| ~~ ~~ + + ~ ~ ~                                 | - +accaa=    | a a a tottoca | tatatittaa | ctcaacttac   | taaaatcgat | 120  |

|   |              |              | cataaaata    | aataaataaq   | taaacaaaat   | 1260 |
|---|--------------|--------------|--------------|--------------|--------------|------|
|   | gcatggctaa   |              |              |              |              |      |
| gaagattgcc                                      | tgctctctct   | gtgcctagcc   | tcaaagcgtt   | catcatacat   | cataccttta   | 1320 |
| agattgctat                                      | attttgggtt   | attttcttga   | caggagaaaa   | agatctaaag   | atcttttatt   | 1380 |
| ttcatctttt                                      | ttggttttct   | tggcatgact   | aagaagctta   | aatgttgata   | aaatatgact   | 1440 |
| agttttgaat                                      | ttacaccaag   | aacttctcaa   | taaaagaaaa   | tcatgaatgc   | tccacaattt   | 1500 |
| caacatacca                                      | caagagaagt   | taatttctta   | acattgtgtt   | ctatgattat   | ttgtaagacc   | 1560 |
| ttcaccaagt                                      | tctgatatct   | tttaaagaca   | tagttcaaaa   | ttgcttttga   | aaatctgtat   | 1620 |
| tcttgaaaat                                      | atccttgttg   | tgtattaggt   | ttttaaatac   | cagctaaagg   | attacctcac   | 1680 |
| tgagtcatca                                      | gtaccctcct   | attcagctcc   | ccaagatgat   | gtgtttttgc   | ttaccctaag   | 1740 |
| agaggttttc                                      | ttcttattt    | tagataattc   | aagtgcttag   | ataaattatg   | ttttctttaa   | 1800 |
| gtgtttatgg                                      | taaactcttt   | taaagaaaat   | ttaatatgtt   | atagctgaat   | ctttttggta   | 1860 |
| actttaaatc                                      | tttatcatag   | actctgtaca   | tatgttcaaa   | ttagctgctt   | gcctgatgtg   | 1920 |
| tgtatcatcg                                      | gtgggatgac   | agaacaaaca   | tatttatgat   | catgaataat   | gtgctttgta   | 1980 |
| aaaagattto                                      | aagttattag   | gaagcatact   | ctgtttttta   | atcatgtata   | atattccatg   | 2040 |
| atacttttat                                      | agaacaattc   | tggcttcagg   | aaagtctaga   | agcaatattt   | cttcaaataa   | 2100 |
| aaggtgttta                                      | aacttt       |              |              |              |              | 2116 |
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| ggaggcggc                                       | g gcgctggtgg | ctgcagctgg   | ggtgaggcgc   | gaggcggcgc   | actcgacggc   | 120  |
| tgactggag                                       | c ageggtaaag | gcgaggatgg   | agaccgaagg   | , atataagtca | aagagtacag   | 180  |
| cagaaaatg                                       | t ctactgaacg | g gacttcttgg | acaagcctgt   | . ccaccattca | . gaaaatagcc | 240  |
| ctgggcctt                                       | g ggateccage | : cagtgcaaca | gttgcctata   | tcctataccg   | caggtatagg   | 300  |
| gaaagcaga                                       | g aagagegget | gacatttgtt   | ggggaagatg   | g acattgagat | agagatgcgg   | 360  |
| gttccccag                                       | g aggctgtgaa | a actcatcatt | ggccggcaag   | g gagccaatat | : taaacagctg | 420  |
| cggaaacag                                       | a caggtgctcg | g gattgatgtg | g gacacagagg | g atgraggcga | tgagcgagtg   | 480  |
| ctgcttatc                                       | a gtggttttc  | c tgttcaggtg | g tgcaaggcca | a aagcagcaat | ccatcagatc   | 540  |
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| acctgtgaca aagaatcaga agggacatta ctactatcaa gacttataaa aatctcagg | ;a /20   |
|--|----------|
| acacagaagg aagtggcagc agccaagcat ttgatactgg agaaagtttc agaagatga | aa 780   |
| gaacttegga agagaattge teattetgea gaaaceaggg teecaegeaa acageeaat | c 840    |
| agtgtgagaa gagaagacat gacagagcca ggtggagctg gagagccagc attatggaa | aa 900   |
| aacaccagtt ctagcatgga geegaetgea eeeetggtga eteeteeace caaaggagg | ga 960   |
| ggcgacatgg ctgtggtagt gtcaaaggaa ggttcctggg agaaacctag tgatgaca  | gc 1020  |
| tttcagaagt ctgaageeca ggccateeca gagatgeeca tgtttgaaag tatgtaac  | aa 1080  |
| agagggagcc cattaatcat tgaagataga aatactggct tagatattgg agagatta  | ga 1140  |
| aggaatgcct totcagtotg agcagocagt ataatatggo tggggtcagg acagaaga  | tt 1200  |
| cggtaagata ttttgaagat gggatggagg tagaatccac accaacagtt gctggaca  | gt 1260  |
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| attgtaggag agaacttggt gootottoca ototgaagca aagttgatga aagtotto  | tt 1380  |
| ccttttccca aaacccaacc tgaactactt ctttcttgga gacagactat attgagac  | aa 1440  |
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| gatatttgta gtttattatt taccccaaac tttctgtgtc tgggtaccct ctgagtag  | ggc 1560 |
| ctataattee tgeetteact gtatgeattt tatgtaaget ageagaeeta tgtggtga  | aga 1620 |
| atgcacagga gcttggaggt ataaatagac agggtgggaa agagagagct cctttcgo  | cca 1680 |
| tgttttacca gcccgctctg ttataacctc ttaggttata ttctttaatt tccaacct  | ttt 1740 |
| taggitagtt totgiaacag aacaaatgag totgggataa agtootoaaa gtaottoa  | aaa 1800 |
| tggtaattgt tttgtttttg taacggetta acaaataace taggttttet gttta     | 1855     |
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| acaccacgat cattcagetg cagecettge aggageeega gagetgggeg egeaege   | aga 12   |
| gtggcctgca gtcctacctg ctccagttcc acggcctcgt gcgcctggtg caccagg   | agc 18   |
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| acacctgtgt             | gcagtatgtg   | cagaaacata | tttccgcgga | aaacacgaaa | gggagccaaa | 480  |
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| gtgtggctgt             | aggcatcttc   | ctgtgcacag | gtggacggcg | atgttaatta | ctctccagcc | 600  |
| ccgtcagaag             | gggctggatt   | gatggaggct | ggcaagggaa | agtttcagct | cactgtgaag | 660  |
| ccagactccc             | caactgaaac   | accagaaggt | ttggagtgac | agctcctttc | ttctcccaca | 720  |
| tctgcccact             | gaagatttga   | gggagggag  | atggagagga | gaggtggaca | aagtacttgg | 780  |
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| aagacagaca             | gaatcacctg   | aggcgttcaa | aagatataac | caaataaaca | agtcatccac | 960  |
| aatcaaaata             | caacattcaa   | tacttccagg | tgtgtcagac | ttgggatggg | acgctgatat | 1020 |
| aatagggtag             | aaagaagtaa   | cacgaagaag | tggtggaaat | gtaaaatcca | agtcatatgg | 1080 |
| cagtgatcaa             | ttattaatca   | attaataata | ttaataaatt | tottatattt | aaggca     | 1136 |
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| <400> 64<br>cgggatagct | gtcctgagcc   | ccaagcctct | tectecectg | ctgcccctct | gcagccattc | 60   |
| gggatgggac             | cccctctggg   | gtgtcagcac | gaaagggcta | acgggagccc | cttccttggc | 120  |
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| tctttgcato             | agcctccatc   | ggcctccaga | ccttcaacca | cagtggcatt | tctgttaaca | 300  |
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| gcctgaagct             | ccttaagaag   | agtccacaac | agctggtggg | agggtggggt | gggcctgggt | 720  |
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| gttctccgca | agcctcctgc | agcgcccgcc | tgccaatgtg | aggctggcac | caggctgcag | 960  |
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| agggcagtgg | gtgggtggcg | ggctagagac | ccttgcctgt | gtccgggacc | ctggcgccgc | 1260 |
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| tcaaaaagaa | aaaaaaaaa  | caaacaaaa  | acataaaaac | aaaaaacaa  | aaagaaaaa  | 1680 |
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<211> 1609 <212> DNA <213> Homo sapien

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| aggggctcaa | ggcggcagtc | cgatagtgga | ggccgctgag | aactgtcacg   | gagetgegte   | 240  |
|------------|------------|------------|------------|--------------|--------------|------|
| tgtacagcga | gcatccctta | tttattcagg | gcgagtgtgt | atttggggcg   | gcgtgcaggg   | 300  |
| ggctgacaaa | gaccggagag | ctcccggtgc | ggccgccggc | ggagcgaaga   | ctggaacccg   | 360  |
| tatgagcgcc | ccccagcgcc | cctgagcgct | cgccgccggt | gcacggcgca   | ccccgcggga   | 420  |
| ggcagggatc | agcaaagccg | tgcgccccga | ggcccgcccc | cgtctccgca   | caaagaccga   | 480  |
| gctggaggat | cttcagaaga | agcctccccc | atacctgcgg | aacctgtcca   | gcgatgatgc   | 540  |
| caatgtcctg | gtgtggcacg | ctctcctcct | acccgaccaa | cctccctacc   | acctgaaagc   | 600  |
| cttcaacctg | cgcatcagct | tcccgccgga | gtatccgttc | aagcctccca   | tgatcaaatt   | 660  |
| cacaaccaag | atctaccacc | ccaacgtgga | cgagaacgga | cagatttgcc   | tgcccatcat   | 720  |
| cagcagtgag | aactggaagc | cttgcaccaa | gacttgccaa | gtcctggagg   | ccctcaatgt   | 780  |
| gctggtgaat | agaccgaata | tcagggagcc | cctgcggatg | gacctcgctg   | acctgctgac   | 840  |
| acagaatccg | gagctgttca | gaaagaatgc | cgaagagttc | acceteegat   | tcggagtgga   | 900  |
| ccggccctcc | taactcatgt | tctgaccctc | tgtgcactgg | atcctcggca   | tagcggacgg   | 960  |
| acacacctca | tggactgagg | ccagagcccc | ctgtggccca | ttccccattc   | atttttccct   | 1020 |
| tcttaggttg | ttagtcatta | gtttgtgtgt | gtgtgtggtg | gagggaaggg   | agctatgagt   | 1080 |
| gtgtgtgttg | tgtatggact | cactcccagg | ttcacctggc | cacaggtgca   | cccttcccac   | 1140 |
| accctttaca | ttccccagag | ccaagggagt | ttaagtitge | agttacaggo   | cagttctcca   | 1200 |
| gctctccatc | ttagagagac | aggtcacctt | gcaggcctgc | ttgcaggaaa   | tgaatccagc   | 1260 |
| agccaactcg | aatcccccta | gggctcaggc | actgagggcc | tggggacagt   | ggagcatatg   | 1320 |
| ggtgggagac | agatggaggg | taccctattt | acaactgagt | cagccaagcc   | actgatggga   | 1330 |
| atatacagat | ttaggtgcta | aaccgtttat | tttccacgga | tgagtcacaa   | . tctgaagaat | 1440 |
| caaacttcca | tcctgaaaat | ctatatgttt | caaaaccact | tgccatcctg   | ttagattgcc   | 1500 |
| agttcctggg | accaggcctc | agactgtgaa | gtatatatco | : tecageatte | agtccagggg   | 1560 |
| gagccacgga | aaccatgttc | ttgcttaagc | cattaaagto | : agagatgaa  |              | 1609 |

<sup>&</sup>lt;210> 66

<sup>&</sup>lt;211> 1414 <212> DNA <213> Homo sapien

<sup>&</sup>lt;400> 66

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<210> 67 <211> 1243 <212> DNA

<213> Homo sapien

<400> 67

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600

660

720

| tgagaactgg aagcettgca ccaagaettg ccaagteetg gaggeeetea atgtgetggt          | 420  |
|--|------|
| gaatagaccg aatatcaggg agcccctgcg gatggacctc gctgacctgc tgacacagaa          | 480  |
| teeggagetg tteagaaaga atgeegaaga gtteaeeete egatteggag tggaeeggee          | 540  |
| ctcctaactc atgttctgac cctctgtgca ctggatcctc ggcatagcgg acggacacac          | 600  |
| ctcatggact gaggccagag ccccctgtgg cccattcccc attcatttt cccttcttag           | 560  |
| gttgttagtc attagtttgt gtgtgtgtgt ggtggaggga agggagctat gagtgtgtgt          | 720  |
| gttgtgtatg gactcactcc caggttcacc tggccacagg tgcacccttc ccacaccctt          | 730  |
| tacattcccc agagccaagg gagtttaagt ttgcagttac aggccagttc tccagctctc          | 840  |
| catcttagag agacaggtca ccttgcaggc ctgcttgcag gaaatgaatc cagcagccaa          | 900  |
| ctcgaatccc cctagggctc aggcactgag ggcctgggga cagtggagca tatgggtggg          | 960  |
| agacagatgg agggtaccct atttacaact gagtcagcca agccactgat gggaatatac          | 1020 |
| agatttaggt gctaaaccgt ttattttcca cggatgagtc acaatctgaa gaatcaaact          | 1080 |
| tccatcctga aaatctatat gtttcaaaac cacttgccat cctgttagat tgccagttcc          | 1140 |
| tgggaccagg cctcagactg tgaagtatat atcctccagc attcagtcca gggggagcca          | 1200 |
| cggaaaccat gttcttgctt aagccattaa agtcagagat gaa                            | 1243 |
| <210> 68<br><211> 1507<br><212> DNA<br><213> Homo sapien                   |      |
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| gctggaggcc acgggtgcca cacactcggt cccgacatga tggcgagcat gcgagtggtg          | 120  |
| aaggagetgg aggatettea gaagaageet eecceatace tgeggaacet gteeagegat          | 180  |
| gatgecaatg teetggtgtg geacgetete etectaceeg aggeagaggt tgeggtgage          | 240  |
| cgagatcacg ccattgcact ccagcctggg cagcaaagtg aaactccatc tcaaaaaaaa          | 300  |
| aaaaaaaaag aagcttggca ccagcatctc cttctgatga ggccctcagg aagcttccac          | 360  |
| tcatggtgga aggcaaaggg gagccacgtg tacagatcac attgcaagag aagaagtgaa          | 420  |
| agagagagag agcgagcagg tgccaggatc ttctaaacaa ccagcttttt cagaccaacc          | 480  |
| tecetaceae etgaaageet teaacetgeg cateagette eegeeggagt ateegtteaa          | 540  |
|  |      |

geoteccatg atcasattca caaccaagat ctaccacccc aacgtggacg agaacggaca

gatttgcctg cccatcatca gcagtgagaa ctggaagcct tgcaccaaga cttgccaagt

cctggaggcc ctcaatgtgc tggtgaatag accgaatatc agggagcccc tgcggatgga

| cctcgctgac                                      | ctgctgacac   | agaatccgga | gctgttcaga | aagaatgccg   | aagagttcac   | 780  |
|---|--------------|------------|------------|--------------|--------------|------|
| cctccgattc                                      | ggagtggacc   | ggecetecta | actcatgttc | tgaccctctg   | tgcactggat   | 840  |
| cctcggcata                                      | gcggacggac   | acacctcatg | gactgaggcc | agagccccct   | gtggcccatt   | 900  |
| ccccattcat                                      | ttttcccttc   | ttaggttgtt | agtcattagt | ttgtgtgtgt   | gtgtggtgga   | 960  |
| gggaagggag                                      | ctatgagtgt   | gtgtgttgtg | tatggactca | ctcccaggtt   | cacctggcca   | 1020 |
| caggtgcacc                                      | cttcccacac   | cctttacatt | ccccagagcc | aagggagttt   | aagtttgcag   | 1080 |
| ttacaggcca                                      | gttctccagc   | tctccatctt | agagagacag | gtcaccttgc   | aggcctgctt   | 1140 |
| gcaggaaatg                                      | aatccagcag   | ccaactcgaa | tccccctagg | gctcaggcac   | tgagggcctg   | 1200 |
| gggacagtgg                                      | agcatatggg   | tgggagacag | atggagggta | ccctatttac   | aactgagtca   | 1260 |
| gccaagccac                                      | tgatgggaat   | atacagattt | aggtgctaaa | ccgtttattt   | tccacggatg   | 1320 |
| agtcacaatc                                      | tgaagaatca   | aacttccatc | ctgaaaatct | atatgtttca   | aaaccacttg   | 1380 |
| ccatcctgtt                                      | agattgccag   | ttcctgggac | caggcctcag | actgtgaagt   | atatatcctc   | 1440 |
| cagcattcag                                      | tccaggggga   | gccacggaaa | ccatgttctt | gcttaagcca   | ttaaagtcag   | 1500 |
| agatgaa   |              |            |            |              |              | 1507 |
| <210> 69<br><211> 127<br><212> DNA<br><213> Hom |              |            |            |              |              |      |
| <400> 69<br>cagcggccgg                          | ccgcggggcc   | tttttgtcct | gagggccaga | gaaatggaga   | agggggtggg   | 60   |
| gggacagcca                                      | cgtggccgca   | ggaggattta | caacattttc | tttcgccatc   | gatgttatcg   | 120  |
| caaaatgtgt                                      | gagagaagcg   | gctgcgcagc | ccggacggga | gcgtgagggt   | gegggeeagg   | 180  |
| taagcagcco                                      | eggeggttte   | geegeataeg | ggactgcggg | gegaeegegg   | gcaccagcca   | 240  |
| cgcgcagcgg                                      | cteegegggg   | teteggeegg | gteegegete | : tgaaggatct | cgagagccat   | 300  |
| ggatggtgca                                      | ggcgggaccc   | tegagetgea | gcatctccgg | tgacccgggg   | ttgccgagga   | 360  |
| ggtggagaco                                      | : agcacaggtg | gtccggcccg | ggegeeteeg | g aatccggggg | g tggtcaagac | 420  |
| ggatccccaa                                      | ggctgaggtc   | ggcagteccg | gggactcgca | a gctgttgago | ctgtggagac   | 480  |
| gcggccccgt                                      | gaccgaggca   | cccttcagca | acccggggg  | agcgttccag   | g agactgaact | 540  |
| tctcaaatca                                      | a ctgcttcaac | agcttttaaa | aatctggac  | c tagttactco | tgtcatctat   | 600  |
| gtgtaaagat                                      | ttagaaaaaa   | aatcccaaac | ccaagggtgg | g gcagccgcga | a ggatgtacag | 660  |
| ctcctggcag                                      | g tgtgcgctca | cccggagacc | tgaggtcaa  | a goccaaatg  | c tgagcacctc | 720  |
|   |              |            |            |              |              |      |

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| ggtagctggg  | gttttcaggt | ttctgcctga | catttaatat | cacaaacatt | gageetgete | 840  |
|---|------------|------------|------------|------------|------------|------|
| ctgtccccca  | ccagccacgc | ggcttctcct | cccaaacata | cccagctcct | teccetgeet | 900  |
| tcctgacggg  | taagaaacat | ccattctttc | caattcccca | gcgttttccc | cctaccggaa | 960  |
| atctgatggg  | cttatgacat | catggctggc | tgctgagcga | tgaagtggat | gccacaaaga | 1020 |
| aatccgacat  | atcagataga | ttctgaaatc | ggtttccctc | cagctgtagt | aacaggcgtg | 1080 |
| aagtcaggag  | aatttgagct | ttgtttaaaa | aataaataaa | taaataaata | aaccataaca | 1140 |
| aagtettgee  | ctgtattaaa | tgcaattttc | ttaaaaacaa | gcaaaccttt | tggacatcat | 1200 |
| tttattttaa  | tagaaatgct | gagttttatg | aaactaaagt | ggctaataaa | tcagacctga | 1260 |
| agctttgtgt  | gagtgttcc  |            |            |            |            | 1279 |
|   | o sapien   |            |            |            |            |      |
| <400> 70<br>cagcggccgg  | ccācāāāācc | tttttgtcct | gagggccaga | gaaatggaga | agggggtggg | 60   |
| gggacagcca  | cgtggccgca | ggaggattta | caacattttc | tttcgccatc | gatgttatcg | 120  |
| caaaatgtgt  | gagagaagcg | getgegeage | ccggacggga | gcgtgagggt | gcgggccagc | 180  |
| gttttccccc  | taccggaaat | ctgatgggct | tatgacatca | tggctggctg | ctgagcgatg | 240  |
| aagtggatgc  | cacaaagaaa | tccgacatat | cagatagatt | ctgaaatcgg | tttccctcca | 300  |
| gctgtagtaa  | caggcgtgaa | gtcaggagaa | tttgagcttt | gtttaaaaaa | taaataaata | 360  |
| aataaataaa  | ccataacaaa | gtcttgccct | gtattaaatg | caattttctt | aaaaacaagc | 420  |
| aaaccttttg  | gacatcattt | tattttaata | gaaatgctga | gttttatgaa | actaaagtgg | 480  |
| ctaataaatc  | agacctgaag | ctttgtgtga | gtgttcc    |            |            | 517  |
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| ctcagacacc  | actttgccat | ccactatccg | gcgggtggtg | gtctttggga | tggtttgcat | 120  |

| teccagatet gtgacecceg gaegacegge gegttetgge agaegtggaa ggaetttgag gtecggaatg geaatgagga caccateaag gaaatgetge gtateegge eagegtgaag gecaegga caccateaag gaaatgetge gtateegge eagegtgaag gecaeggaa eagettetgg eagetgagg gecaeggage eagetgagg caactteatg geetegaagga getegaggagg eteggaggaggaggaacaggaacaggaacaggaacaggagaagaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaagaagaagaagaagaagaagaagaaggaaggaaggaaggaaggaaggaagaagaagaagaagaagaagaaggaagaagaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaaggaagaagaagaagaagaagaaggaagaagaaggaagaagaaggaagaagaaggaagaaggaagaagaagaagaagaagaagaaggaagaagaaggaagaagaaggaag |                          |
|--|--------------------------|
| gccacgtaca acacgcaggt caacttcatg gcatcgaga tgctcaaggt ctcgggaggt gccacggtaca acacgcaggt caacttcatg gcctcgaga tgctcaaggt ctcgggagt gccacggaca ccgtgtctga cctggccct gggcagagtg gcatcgacgac cttgggcgcc cagagcagaca gcggcagaga gctgccgcc gggcagagtg gtgaccagcc cttgcgcgcc cagagcaaga tcctgttcgt gaggagtgac gcctcccggg aggactggc aggactggc agagctggc cagcaggtca accccgagga gatccagct ggcgaggaca aggactggc agagctggc ctgaacgag accccgaga gatccagct gctggagcac acgaggtca accccgaga gatccagct ggcgaggacac acgaggtca accccgaga gatccagct ggcgaggacac aggacgaga cagaatggac ctgaacgaag actgacccgt ccctccccca tcccccccc ccacccctc cccaatacag ctacgtttgt acatca  <2210 > 72 <211 > 3041 <212 > DNA <213 > Homo sapien  <400 > 72 aatgatagat cactataggg gaatgggtcc tctaatgctg ctcgagcgc gcagtgtgat ggatcgtggt caccacacac ccacaaacca actctggaat acagtcctt tccccatac ccccatacac ccccaaacca actctggaat acagtcctt tcccacagaa ccagcatgta gagcagagc cacactacac ccacaaacca actctggaat acagtcatt tcccacagaa cttgcccagaacgcc tctgagggt tcagaggtga tacaagtat ctcacacagaa tttgccca aaccaaagct tagttccct tcttggtgat caaagggaat ctgagaataa tgctaatcca gttttgccca aaccaaagct tagttccct tcttggtgat caaagggaat ctgagaataat tgctaatccc atgcagactt cccccagtgt tcagcagcaa aatgatgct acttgcacag cttcagcatg aggccatgga ggcactggg gggacacgaga tggcctctga ctcttcaaac acttcactgc cattctcaaa cattgggaaat ccaatgaaca ccacacagtt agggaaatca ctttttcagt ggcaggtgga gcaggaagaa agcaaattgg caaatattc ccaaagaccag tttcttcaa aggatgcaga tggtgacacg ttccttcaaa ttgctgttc ccaaaggaga agggcacttt cctatgttc tgcaagaaag atgaatgcac ttccttcaaa ttccattgt ggatataaa gagcacaatg gacagaggc ctttcaggt gacagggg ccaaggt gacacaggt gcagtggct ccaatcagca tccattttc ggaacacag gacagagga acgagacacacag acgacagga aggacacacc ctgcatggt gacacagg gcacaggt gacacaggt gcagtggct caaaccacag actgctggg aagaacacct ctgcatggg gacacacgg ggacacagg gacacacaca accacaaga acgacacac ctgcatggg gacacacac ctgcatggg gacacacac ctgcatggg gacacacac ctgcatggg gacacacac ctgcatggg gacacacaca ggacacacac accacacag actgctggga agaacacct ctgcatgtg gtgcctgagaa gggcacacacc caggtgctc aggcacacacac acacacacac | gecateta cagettetge 180  |
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| catteteaaa catgggaaat ecaatgaaca ecacacagtt agggaaatea etttteagt ggcaggtgga gcaggaagaa agcaaattgg caaatattte ecaaggaaga tttettteaa aggatgcaga tggtgacacg tteetteata ttgetgttge ecaagggaga agggcaettt ectatgttet tgcaagaaag atgaatgcae tteacatget ggatattaaa gagcacaatg gacagagtge ettteaggtg geagtggetg ecaateagea teteattgtg eaggatetgg tgaacategg ggcacaggtg aacaceacag actgetgggg aagaacacet etgeatggt gtgetgagaa gggccaetee eaggtgette aggegattea gaagggagca gtgggaagta ateagtttgt ggatettgag gcaactaact atgatggett gaeteeeett eactgtgeag  | tcagcatg atgcccagca 360  |
| ggcaggtgga gcaggaagaa agcaaattgg caaatattte ccaagaccag tttetteaa aggatgcaga tggtgacacg tteetteata ttgetgttge ccaagggaga agggcaettt cetatgttet tgcaagaaag atgaatgcae tteacatget ggatattaaa gagcacaatg gacagagtge ettteaggtg geagtggetg ecaateagea teteattgtg caaggatetgg tgaacategg ggcacaggtg aacaccacag actgetgggg aagaacacet etgeatgtg gtgetgagaa gggccaetee eaggtgette aggegattea gaagggagca gtgggaagta atcagttgt ggatettgag gcaactaact atgatggeet gaeteeeett caetgtgeag  | cttcaaac acttcactgc 420  |
| aggatgcaga tggtgacacg ttccttcata ttgctgttgc ccaagggaga agggcacttt cctatgttct tgcaagaaag atgaatgcac ttcacatgct ggatattaaa gagcacaatg gacagagtgc ctttcaggtg gcagtggctg ccaatcagca tctcattgtg caggatctgg tgaacatcgg ggcacaggtg aacaccacag actgctgggg aagaacacct ctgcatgtgt gtgctgagaa gggccactcc caggtgcttc aggcgattca gaagggagca gtgggaagta atcagtttgt ggatcttgag gcaactaact atgatggcct gactcccctt cactgtgcag  | ggaaatca ctttttcagt 480  |
| cctatgttct tgcaagaag atgaatgcac ttcacatgct ggatattaaa gagcacaatg gacagagtgc ctttcaggtg gcagtggctg ccaatcagca tctcattgtg caggatctgg tgaacatcgg ggcacaggtg aacaccacag actgctgggg aagaacacct ctgcatgtgt gtgctgagaa gggccactcc caggtgcttc aggcgattca gaagggagca gtgggaagta atcagtttgt ggatcttgag gcaactaact atgatggcct gactcccctt cactgtgcag   | aagaccag tttctttcaa 540  |
| gacagagtge ettteaggtg geagtggetg ecaateagea teteattgtg eaggatetgg tgaacategg ggeacaggtg aacaceacag actgetgggg aagaacacet etgeatgtg gtgetgagaa gggeeactee eaggtgette aggegattea gaagggagea gtgggaagta ateagtttgt ggatettgag geaactaact atgatggeet gaeteeeett eactgtgeag   | aagggaga agggcacttt 600  |
| tgaacatcgg ggcacaggtg aacaccacag actgctgggg aagaacacct ctgcatgtgt gtgctgagaa gggccactcc caggtgcttc aggcgattca gaagggagca gtgggaagta atcagtttgt ggatcttgag gcaactaact atgatggcct gactcccctt cactgtgcag  | atattaaa gagcacaatg 660  |
| gtgctgagaa gggccactcc caggtgcttc aggcgattca gaagggagca gtgggaagta atcagtttgt ggatcttgag gcaactaact atgatggcct gactcccctt cactgtgcag  | tcattgtg caggatctgg 720  |
| atcagitigt ggatettgag gcaactaact atgatggeet gaeteeeett caetgtgeag  | gaacacct ctgcatgtgt 780  |
|  | agggagca gtgggaagta 840  |
|  | actcccctt cactgtgcag 900 |
| tcatagecca caatgetgtg gtecatgaae tecagagaaa teaacageet catteacetg  | caacagcct cattcacctg 960 |

| aagtto | cagga | gcttttactg | aagaataaga | gtctggttga | taccattaag | tgcctaattc | 1020 |
|--------|-------|------------|------------|------------|------------|------------|------|
| aaatg  | ggagc | agcggtggaa | gcgaaggatc | gcaaaagtgg | ccgcacagcc | ctgcatttgg | 1080 |
| cagct  | gaaga | agcaaatctg | gaactcattc | gcctctttt  | ggagetgeee | agttgcctgt | 1140 |
| ctttt  | gtgaa | tgcaaaggct | tacaatggca | acactgccct | ccatgttgct | gccagcctgc | 1200 |
| agtat  | cggtt | gacacaatta | gatgctgtcc | gcctgttgat | gaggaaggga | gcagacccaa | 1260 |
| gtact  | cggaa | cttggagaac | gaacagccag | tgcatttggt | tecegatgge | cctgtgggag | 1320 |
| aacag  | atccg | acgtatcctg | aagggaaagt | ccattcagca | gagagctcca | ccgtattagc | 1380 |
| tccat  | tagct | tggagcctgg | ctagcaacac | tcactgtcag | ttaggcagtc | ctgatgtatc | 1440 |
| tgtac  | ataga | ccatttgcct | tatattggca | aatgtaagtt | gtttctatga | aacaaacata | 1500 |
| tttag  | ttcac | tattatatag | tgggttatat | taaaagaaaa | gaagaaaaat | atctaatttc | 1560 |
| tettg  | gcaga | tttgcatatt | tcatacccag | gtatctggga | tctagacatc | tgaatttgat | 1620 |
| ctcaa  | tggta | acattgcctt | caattaacag | tagcttttga | gtaggaaagg | actttgattt | 1630 |
| gtggc  | acaaa | acattattaa | tatagctatt | gacagtttca | aagcaggtaa | attgtaaatg | 1740 |
| tttct  | ttaag | aaaaagcatg | tgaaaggaaa | aaggtaaata | cagcattgag | gcttcatttg | 1800 |
| gcctt  | agtcc | ctgggagtta | ctggcgttgg | acaggettea | gtcattggac | tagatgaaag | 1860 |
| gtgtc  | catgg | ttagaatttg | atctttgcaa | actgtatata | attgttattt | ttgtccttaa | 1920 |
| aaata  | ttgta | catacttggt | tgttaacatg | gtcatatttg | aaatgtataa | gtccataaaa | 1980 |
| tagaa  | aagaa | caagtgaatt | gttgctattt | aaaaaaattt | tacaattctt | actaaggagt | 2040 |
| tttta  | ttgtg | taatcactaa | gtctttgtag | ataaagcaga | tggggagtta | cggagttgtt | 2100 |
| ccttt  | actgg | ctgaaagata | tattcgaatt | gtaaagatgc | tttttctcat | gcattgaaat | 2160 |
| tatac  | attat | ttgtagggaa | ttgcatgctt | tttttttt   | ttctcccgag | acagggtctt | 2220 |
| gctct  | ggcgc | ccaggctgga | gtacagtggc | atgatcttgg | ctcacttcag | ccttgacttg | 2280 |
| ggctc  | aagtg | atcctcctac | ctgagecttc | tgagtaactg | ggactacagg | tgtgcactcc | 2340 |
| tegee  | tggct | aatttttat  | tttttgtaca | ggcaggatct | tgccaccttg | cccaggctgg | 2400 |
| tcttg  | aactc | ctgagctcat | gccatctgcc | tgccttagtc | tcccaaaatg | ctgggattac | 2460 |
| aggag  | tgagc | caccatgeec | ggctggcagt | tgcatggaag | agaacacctc | tttatggctt | 2520 |
| accct  | ctaga | atttctaatt | tatgtgttct | gttgaaattt | ttgtttttt  | acctttattg | 2580 |
| aaaca  | acaaa | aagtcagtat | tgaaacatat | cttcctgttt | tctgttgtca | aatgatgata | 2640 |
| atgtg  | ccatg | atgttttata | tatatcatto | agaaaaagtt | ttattttta  | ataacattct | 2700 |
| attaa  | catta | ttttgcttgc | cgctggcatg | cctgaggaat | gtatttggct | ttgattacac | 2760 |
| actaa  | gttt  | tgtaataaat | ttgactcatt | aaaaaccttt | ttttttaaa  | aaaaaaaaa  | 2820 |

| agaaaatctc attagtgaac  | ttatctttgc | agctgagtac | ttaaattott | tttaaaaaga | 2880 |
|--|------------|------------|------------|------------|------|
| taccctttgg attgatcaca  | ttgtttgacc | cagtatgtct | tgtagacacg | ttagttataa | 2940 |
| tcaccttgta tctctaaata  | tggtgtgata | tgaaccagtc | cattcacatt | ggaaaaactg | 3000 |
| atggttttaa ataaactaat  | tcactaatat | tatttgtctt | a          |            | 3041 |
| <210> 73<br><211> 1193<br><212> DNA<br><213> Homo sapien<br><400> 73 |            |            |            |            |      |
| acatccagct gcctgagacc  | ctcctgcagc | cttctcaagg | gacageeeca | ctctgcctct | 60   |
| tgctcctcca gggcagcacc  | atgcagcccc | tgtggctctg | ctgggcactc | tgggtgttgc | 120  |
| ccctggccag ccccggggcc  | gccctgaccg | gggagcagct | cctgggcagc | ctgctgcggc | 180  |
| agetgeaget caaagaggtg  | cccaccctgg | acagggccga | catggaggag | ctggtcatcc | 240  |
| ccacccacgt gagggcccag  | tacgtggccc | tgctgcagcg | cagccacggg | gaccgctccc | 300  |
| gcggaaagag gttcagccag  | agcttccgag | aggtggccgg | caggttcctg | gcgttggagg | 360  |
| ccagcacaca cctgctggtg  | ttcggcatgg | agcagcggct | gccgcccaac | agcgagctgg | 420  |
| tgcaggccgt gctgcggctc  | ttccaggaat | gtacattgac | ctgcagggga | tgaagtgggc | 480  |
| cgagaactgg gtgctggagc  | ccccgggctt | cctggcttat | gagtgtgtgg | gcacctgccg | 540  |
| geageeeeeg gaggeeetgg  | ccttcaagtg | gccgtttctg | gggcctcgac | agtgcatcgc | 600  |
| ctcggagact gactcgctgc  | ccatgatcgt | cagcatcaag | gagggaggca | ggaccaggcc | 660  |
| ccaggtggtc agcctgccca  | acatgagggt | gcagaagtgc | agctgtgcct | cggatggtgc | 720  |
| gctcgtgcca aggaggctcc  | agccataggc | gcctagtgta | gccatcgagg | gacttgactt | 780  |
| gtgtgtgttt ctgaagtgtt  | cgagggtacc | aggagagctg | gcgatgactg | aactgctgat | 840  |
| ggacaaatge tetgtgetet  | ctagtgagcc | ctgaatttgc | ttcctctgac | aagttacctc | 900  |
| acctaatttt tgcttctcag  | gaatgagaat | ctttggccac | tggagagccc | ttgctcagtt | 960  |
| ttctctattc ttattattca  | ctgcactata | ttctaagcac | ttacatgtgg | agatactgta | 1020 |
| acctgagggc agaaagccca  | atgtgtcatt | gtttacttgt | cctgtcactg | gatctgggct | 1080 |
| aaagtcctcc accaccactc  | tggacctaag | acctggggtt | aagtgtgggt | tgtgcatccc | 1140 |
| caatccagat aataaagact  | ttgtaaaaca | tgaataaaac | acattttatt | cta        | 1193 |

<sup>&</sup>lt;210> 74

<sup>&</sup>lt;211> 1725

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo sapien

| ttttttttca | cctaaattgt | ttttacttgc | ttcaccttaa | atcagttatt | cggttccttt | 60   |
|------------|------------|------------|------------|------------|------------|------|
| gaagaacatg | tagctttagc | aacccccaaa | gtgaacccag | atacccagaa | ctactgattg | 120  |
| ttacatcaga | gtttttgaaa | acgccaatac | tggaaacagg | ccgacagtta | aggagaggca | 180  |
| ggcctcaaac | taaagcagaa | ccttctcaaa | actacttata | tttcaagaga | tggattaagc | 240  |
| gaaaagagct | gtacaaagaa | ataagccatt | ttctatgcgc | ctgttatatg | caaggcactg | 300  |
| cactcccttg | tcacttgtta | atttacaaga | agactcgccg | gtccagggaa | gggagagggg | 360  |
| aataggtgca | aggtttgtgg | agggcggcaa | actgcactcc | atgtggcgcg | aggctgcgct | 420  |
| tgcggagatc | cgggttctgg | tgcctggcgc | cggcagccag | taggggcgcg | cgtcctgcgc | 480  |
| agttgccgcg | cgcggcctcc | tcgggcttca | ggccgggcca | atcaggaggc | gggcgggagg | 540  |
| gggcgcggta | ccaggaggga | acgggacggg | tegggetteg | aggtcagctg | gtccgcaggg | 600  |
| aagcctttgg | cttccccacc | ggcaatggag | cggttgacgt | tgcctctcgg | cggcgcggcg | 660  |
| gcggtggacg | agtacctgga | gtaccggagg | tataaacaac | ataaaactga | cttggaagcg | 720  |
| attcctcagc | agtgccccat | tgatctgccc | tgccaagtga | ctggctgcca | gtgcagggct | 780  |
| tacctttatg | tccccttgaa | tggtagccag | cccattcgct | gcaggtgcaa | acactttgct | 840  |
| gatcagcaca | gtgctgcgcc | tggctttaca | tgcaatacat | gttccaagtg | ttcaggattc | 900  |
| catagctgct | tcacttgtgc | ttgtggtcag | cctgcatatg | cccatgacac | agtagtggaa | 960  |
| actaagcaag | aaagattggc | tcaggaaaaa | ccagtgggac | aggacattcc | ttatgcagcc | 1020 |
| atgggaggat | taactggttt | cagctcgctg | gcggaaggct | acatgcggtt | agatgacagt | 1030 |
| gggattgtag | gtacaagtag | tcaagtttct | tcattaagga | gacctgaaga | ggatgatatg | 1140 |
| gctttctttg | aaagacgata | ccaggaaagg | atgaaaatgg | aaaaggctgc | taagtggaaa | 1200 |
| ggaaaagctc | cattgccatc | agctacaaaa | ccttcatgaa | gactattgga | gaaattaaaa | 1260 |
| ccatcatcca | agtatctttt | tcatgtttat | ttaaatgtaa | taatacagtt | tatttttcct | 1320 |
| gaaattattt | acttttttt  | tttactgtat | aaatgtcttt | tgggatgttt | ccttaattta | 1380 |
| tttaaataac | taaaaatgtc | tattactttt | gtcaaaactc | ataatttact | actttgtatg | 1440 |
| tacctttctt | tctcctgaca | aatgaggtta | ttttatatga | gtctgtctga | gagtacagta | 1500 |
| aatgttttta | gtacataata | atttaactgt | ttcaggtatt | taaaaaatta | aagatattat | 1560 |
| caagggtttt | ggacaaacat | atgagccatt | tttttgtcat | tcaaaatagt | aagttaaaaa | 1620 |
| caagagaaca | aaagaaaaaa | atagttaaaa | tcattaattt | ttttatttt  | caaactttgt | 1630 |
| aatgttatgt | tttcaaataa | aaactatcto | aaaattttac | aacca      |            | 1725 |

| <210> /5   |           |            |            |            |              |      |
|--|-----------|------------|------------|------------|--------------|------|
| <211> 670  |           |            |            |            |              |      |
| <212> DNA<br><213> Homo                          | sapien    |            |            |            |              |      |
| (213) HOMO                                       | sabien    |            |            |            |              |      |
| <400> 75   |           |            |            |            |              |      |
| ggaggttcag g                                     | aagccattt | tctttcgcgt | ctgcggtgct | cggagtgtgg | tacttctcct   | 60   |
| agttgcagtc a                                     | ggcttcata | cgctattgtc | ctgcccgtta | gagcagccag | cgggtacaga   | 120  |
| atggattttg g                                     | aagagggag | tcaccactgg | acctccaagg | aagccacgtg | cagacatcta   | 180  |
| caaccttcga t                                     | ctcctgacg | agtttattgt | tggccaaaac | caggctttga | ttgaaccagg . | 240  |
| atgaatgcgg g                                     | tgttggaag | tagaatatat | atatacatat | aaaattggtt | gggagccacg   | 300  |
| tgtaccagtg t                                     | gtgttgatc | ttggcttgat | tcagtctgcc | ttgtaacaga | aactggcgat   | 360  |
| ggaatatgag a                                     | ggagccctc | tggaaagaaa | aggacagacc | ctgtgctttc | atgaaagtga   | 420  |
| agatctggct g                                     | aaccagttc | cacaaggtta | ctgtatacat | agcctgagtt | taaaaggctg   | 480  |
| tgcccacttc a                                     | agaatgtca | ttgttagact | ttgaaatttc | taactgccta | cctgcataaa   | 540  |
| gaaaataaaa t                                     | cttttaaat | Caaaaaaaaa | acaaaacaa  | aaaaacttgt | ggtgggggc    | 600  |
| cttgggcccc g                                     | gaaaaggtt | ttttaaccct | tccgttgggg | 99999999cc | ccttggtttg   | 660  |
| eggeeeetg  |           |            |            |            |              | 670  |
| <210> 76<br><211> 964<br><212> DNA<br><213> Homo | sapien    |            |            |            |              |      |
| <400> 76<br>ggaggttcag g                         | aagccattt | tctttcgcgt | ctgcggtgct | cggagtgtgg | tacttctcct   | 60   |
| agttgcagtc a                                     | ggcttcata | cgctattgtc | ctgcccgtta | gagcagccag | cgggtacaga   | 120  |
| atggattttg g                                     | aagagggag | tcaccactgg | acctccaagg | aagccacgtg | cagacatcta   | 180  |
| caaccttcga t                                     | ctcctgacg | agtttattgt | tggccaaaac | caggctttga | ttgaaccagg   | 240  |
| atgaatgcgg g                                     | tgttggaag | tagaatatat | atatacatat | aaaattgaaa | ctggcgatgg   | 300  |
| aatatgagag g                                     | agccctctg | gaaagaaaag | gacagaccct | gtgctttcat | gaaagtgaag   | 360  |
| atctggctga a                                     | ccagttcca | caaggttact | gtatacatag | cctgagttta | aaaggctgtg   | 420  |
| cccacttcaa g                                     | aatgtcatt | gttagacttt | gaaatttcta | actgcctacc | tgcataaaga   | 480  |
|  |           |            |            |            |              | - 40 |
| aaataaaatc t                                     | tttaaatca | aaaaaaaaa  | aaaggggggg | cegecetttt | ttttttttt    | 540  |
| tttgggaggt c                                     |           |            |            |            |              | 600  |
|  | cccaacaat | tcttgggtaa | ttctttcccc | aaaaggtaaa | tatacccgaa   |      |

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|  |            | 120        |            |            |     |
|--|------------|------------|------------|------------|-----|
| tecgeccaac gacaagcaga  | aacaaaagga | gggcggcgag | ccaacacatc | agcaaggcga | 780 |
| caagacccac agccaagagg  | caggcggccg | acagtgacat | aacaacgagc | ggccccacgg | 840 |
| caacgacaac cgacgataag  | aacgacgtgt | gtgaagacac | accccaccgc | cgaacgacgg | 900 |
| gatggcacca gcgagaccta  | tagtacgcga | agcagccaag | acgaagagcg | gggaagcaag | 960 |
| gaaa   |            |            |            |            | 964 |
| <210> 77 <211> 269 <212> DNA <213> Homo sapien   |            |            |            |            |     |
| <400> 77 cctctctggc ggcggctgct   | gctagggagt | cgacttctcc | tgccttgtaa | cagaaactgg | 60  |
| cgatggaata tgagaggagc  | cctctggaaa | gaaaaggaca | gaccctgtgc | tttcatgaaa | 120 |
| gtgaagatct ggctgaacca  | gttccacaag | gttactgtat | acatageetg | agtttaaaag | 180 |
| gctgtgccca cttcaagaat  | gtcattgtta | gactttgaaa | tttctaactg | cctacctgca | 240 |
| taaagaaaat aaaatctttt  | aaatcaaaa  |            |            |            | 269 |
| <210> 78 <211> 928 <212> DNA <213> Homo sapien <220> <221> misc_feature <222> (855)(855) <223> n=a, c, g or t <400> 78 |            |            |            |            |     |
| ggaggttcag gaagccattt  | tetttegegt | ctgcggtgct | cggagtgtgg | tacttctcct | 60  |
| agttgcagtc aggcttcata  | cgctattgtc | ctgcccgtta | gagcagccag | cgggtacaga | 120 |
| atggattttg gaagagggag  | tcaccactgg | acctccaagg | aagccacgtg | cagacatcta | 180 |
| caacettega teteetgaeg  | agtttattgt | tggccaaaac | caggetttga | ttgaaccagg | 240 |
| atgaatgcgg gtgttggaag  | tagaatatat | atatacatat | aaaattggtt | gggagccacg | 300 |
| tgtaccagtg tgtgttgatc  | ttggcttgat | tcagtctgcc | ttgtaacaga | aactggcgat | 360 |
| ggaatatgag aggagccctc  | tggaaagaaa | aggacagacc | ctgtgctttc | atgaaagtga | 420 |
| agatctggct gaaccagttc  | cacaaggtta | ctgtatacat | agcctgagtt | taaaaggctg | 480 |
| tgcccacttc aagaatgtca  | ttgttagact | ttgaaatttc | taactgccta | cctgcataaa | 540 |
| gaaaataaaa tottttaaat  | caaaaaaaa  | aaccaaaaa  | aatatctata | aacacaaaga | 600 |
| aaaacaaatt caaacttaca  | cattatccaa | tcattctatt | attataaata | tatgaaaaaa | 660 |

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| aaaaaaactt g                             | ggggcccgc t | tggcccagg  | gggttttaaa  | tctggaccaa   | tttattcttg   | 720  |
|--|-------------|------------|-------------|--------------|--------------|------|
| gggcccaatt t                             | ggggctcat ( | ttggccaat  | tgccccttta  | ggggaggcgt   | ttacaattca   | 780  |
| tgggccggtt t                             | ttcaaacgg   | tggtatttgg | aaaacccctg  | gggttaccca   | atttttaaac   | 840  |
| ccttgggaaa a                             | aatnccctt   | ttttcacagt | gggggaaat   | tccgaaggcc   | ccgccccgtg   | 900  |
| aggccttttc c                             |             |            |             |              |              | 928  |
| <210> 79 <211> 2448 <212> DNA <213> Homo | sapien      |            |             |              |              |      |
| <400> 79                                 | tcactagtag  | cattgttaat | attaaacaag  | atagctaata   | gctttaaagc   | 60   |
| gtctaatcac                               |             |            |             |              |              | 120  |
| aaacagcgct                               |             |            |             |              |              | 180  |
| gggatcagcg                               |             |            |             |              |              | 240  |
| ccgctgcaag                               |             |            |             |              |              | 300  |
| acaaaatctt                               |             |            |             |              |              | 360  |
|  |             |            |             |              | : tcagaggacg | 420  |
|  |             |            |             |              | tcagctactc   | 480  |
|  |             |            |             |              | g gggcgtcctg | 540  |
|  |             |            |             |              | a gaagcaggaa | 600  |
|  |             |            |             |              | a gccatatgga | 660  |
|  |             |            |             |              | a tgggtggtga | 720  |
|  |             |            |             |              | g gcgaagggtc | 780  |
|  |             |            |             |              | t ctgtccctga | 840  |
|  |             |            |             |              | c gttcgctggt | 900  |
|  |             |            |             |              | g tegeegegga | 960  |
|  |             |            |             |              |              | 1020 |
|  |             |            |             |              | g cagataaagt | 1080 |
|  |             |            |             |              | t gtagaatatt | 1140 |
|  |             |            |             |              | a ttgggtttag | 1200 |
|  |             |            |             |              | c caggaagcag | 126  |
| ctagtcttt                                |             |            |             |              | a gccttcttt  | 132  |
|  |             | ~ ~~~~~~~~ | а паасослов | ia Luulututt | _ ~~~~~~~~~~ |      |

| tggaaggtgt                                   | gcatgtggaa   | gaggaagaag   | gagaaaaaac   | agaagatgaa  | tjetetggtag  | 1380 |
|--|--------------|--------------|--------------|-------------|--------------|------|
| aaaataatga                                   | taacatagat   | gaaactgaag   | gctcggaaga   | ggatgataaa  | gaaaatgata   | 1440 |
| agactgaaga                                   | aatgccaaat   | gattcagtcc   | ttgaaaacaa   | gtctcttcaa  | gaaaatgagg   | 1500 |
| aggaggagat                                   | tgggaaccta   | gagettgeet   | gggatatgct   | ggatttagca  | aagatcattt   | 1560 |
| ttaaaaggca                                   | agaaacaaaa   | gaagcacagc   | tttatgctgc   | ccaggcacat  | cttaaactcg   | 1620 |
| gagaagttag                                   | tgttgaatct   | gaaaactatg   | tgcaagctgt   | ggaggagttc  | cagtcctgcc   | 1680 |
| ttaacctgca                                   | ggaacagtac   | ctggaagccc   | acgaccgtct   | ccttgcagag  | acccactacc   | 1740 |
| agctgggctt                                   | ggcttatggg   | tacaactctc   | agtatgatga   | ggcagtggca  | cagttcagca   | 1300 |
| aatctattga                                   | agtcattgag   | aacagaatgg   | ctgtactaaa   | cgagcaggtg  | aaggaggctg   | 1860 |
| aaggatcgto                                   | : tgctgaatac | aagaaagaaa   | ttgaggaact   | aaaggaactg  | ctacccgaaa   | 1920 |
| ttagagagaa                                   | gatagaagat   | gcaaaggagt   | ctcagcgtag   | tgggaatgta  | gctgaactgg   | 1980 |
| ctctgaaagc                                   | tactctggtg   | gagagttcta   | cttcaggttt   | cactcctggt  | ggaggaggct   | 2040 |
| cttcagtctc                                   | catgattgcc   | agtagaaagc   | caacagacgg   | tgcttcctca  | tcaaattgtg   | 2100 |
| tgactgatat                                   | ttcccacctt   | gtcagaaaga   | agaggaaacc   | agaggaagag  | agtccccgga   | 2160 |
| aagatgatgo                                   | aaagaaagcc   | aaacaagagc   | cggaggtgaa   | cggaggcagt  | ggggatgctg   | 2220 |
| tccccagtg                                    | g aaatgaagtt | tcggaaaaca   | tggaggagga   | ggctgagaat  | : caggctgaaa | 2280 |
| gccgggcag                                    | agtggagggg   | acagtggagg   | ctggagctac   | agttgaaago  | actgcatgtt   | 2340 |
| aagaggggg                                    | c acageceted | : tcccaaggga | aagtgttttt   | gtatataatg  | g tattttttca | 2400 |
| cttttggag                                    | g attotttte  | ; tataacttca | ataaagattg   | taagcaaa    |              | 2448 |
| <210> 80 <211> 16 <212> DN <213> Ho <400> 80 | A            |              |              |             |              |      |
| tgacgcgaa                                    | t gacgctgcgd | c cagtcaggco | gcagccccgc   | tgcttggcc   | c gtcgggcccg | 60   |
| ccttggccg                                    | g ctcggcccg  | c ccccggccct | ccctgcacgg   | g cctcccgtg | c gcccctgtca | 120  |
| gactgtggc                                    | g gccggtcgc  | g eggtgegete | c teecteect  | g cccgcagcc | t ggagaggcgc | 180  |
| ttcgtgctg                                    | c acacccccg  | c gttcctgcc  | g gcaccgcgc  | tgccctctg   | c cgcgctccgc | 240  |
| cctgccgcc                                    | g accgcacgc  | c cgccgcggg  | a catggcacao | e gcaccggca | c gctgccccag | 300  |
| cgcccgggg                                    | c teeggggae  | g gcgagatgg  | g caageeeag  | g aacgtggcg | c tcatcaccgg | 360  |
| tatcacago                                    | c caggatggt  | t cctacctgg  | c tgagttcct  | g ctggagaaa | g gctatgaggt | 420  |
|  |              |              |              |             |              |      |

ccatggaatt gtacggcggt ccagttcatt taatacgggt cgaattgagc atctgtataa 480

| gaatccccag | gctcacattg   | aaggaaacat | gaagttgcac | tatggcgatc  | tcactgacag | 540  |
|------------|--------------|------------|------------|-------------|------------|------|
| tacctgcctt | gtgaagatca   | ttaatgaagt | aaagcccaca | gagatctaca  | accttggagc | 600  |
| ccagagecac | gtcaaaattt   | cctttgacct | cgctgagtac | actgcggacg  | ttgacggagt | 660  |
| tggcactcta | cgacttctag   | atgcagttaa | gacttgtggc | cttatcaact  | ctgtgaagtt | 720  |
| ctaccaagcc | tcaacaagtg   | aactttatgg | gaaagtgcag | gaaatacccc  | agaaggagac | 780  |
| cacccctttc | tatccccggt   | caccctatgg | agctaatttc | gttactcgaa  | aaattagccg | 840  |
| gtcagtagct | aagatttacc   | ttggacaact | ggaatgtttc | agtttgggaa  | atctggatgc | 900  |
| caaacgagat | tggggccatg   | ccaaggacta | tgtggaggct | atgtggttga  | tgttgcagaa | 960  |
| tgatgagccg | gaggacttcg   | ttatagctac | tggggaggtc | catagtgtcc  | gggaatttgt | 1020 |
| cgagaaatca | ttcttgcaca   | ttggaaaaac | cattgtgtgg | gaaggaaaga  | atgaaaatga | 1080 |
| agtgggcaga | tgtaaagaga   | ccggcaaagt | tcacgtgact | gtggatctca  | agtactaccg | 1140 |
| gccaactgaa | gtggactttc   | tgcagggcga | ctgcaccaaa | gcgaaacaga  | agctgaactg | 1200 |
| gaagccccgg | gtcgctttcg   | atgagctggt | gagggagatg | gtgcacgccg  | acgtggagct | 1260 |
| catgaggaca | aaccccaatg   | cctgagcagc | geeteggage | ceggecegee  | ctccggctac | 1320 |
| aatccccgca | gagtctccgg   | tgcagacgcg | ctgcggggat | ggggagcggc  | gtgccaatct | 1380 |
| gcgggtccc  | tgeggeeect   | gctgccgctg | cgctgtcccg | gccgcaagag  | caaaaccacc | 144  |
| ccgccgaggt | : tigtagcagc | cgggatgtga | ccctccaggg | tttgggtcgc  | titgcgtttg | 1500 |
| tcgaagcct  | c ctctgaatgg | ctttgtgaas | tcaagatgtt | ttaatcacat  | teactttact | 156  |
| tgaaattatg | g ttgttacaca | acaaattgtg | gggccttcaa | a attgttttt | tcttttcata | 162  |
| ttaaaaatg  | tctttctgtg   | aactagca   |            |             |            | 164  |

<210> 81

<211> 2595

<212> DNA

<213> Homo sapien

<400> 81
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gactgtggcg gccggtcgcg cggtgcgct tccctccctg cccgcagcct ggagaggcgc 180

ttcgtgctgc acacccccgc gttcctgccg gcaccgcgcc tgccctctgc cgcgctccgc 240

cctgccgccg accgcacgc cgccgcggga catggcacac gcaccggcac gctgcccag 300

cgcccggggc tccggggacg gcgagatggg caagcccagg aacgtggcgc tcatcaccgg 360

tatcacaggc caggatggtt cctacctggc tgagttcctg ctggagaaag gctatgaggt 420

| catggaatt gt  | tacggcggt  | ccagttcatt   | taatacgggt   | cgaattgagc | atctgtataa   | 480  |
|---------------|------------|--------------|--------------|------------|--------------|------|
| gaateeceag ge | ctcacattg  | aaggaaacat   | gaagttgcac.  | tatggcgatc | tcactgacag   | 540  |
| tacctgcctt g  | tgaagatca  | ttaatgaagt   | aaagcccaca   | gagatetaca | accttggagc   | 600  |
| ccagagccac g  | tcaaaattt  | cctttgacct   | cgctgagtac   | actgcggacg | ttgacggagt   | 660  |
| tggcactcta c  | gacttctag  | atgcagttaa   | gacttgtggc   | cttatcaact | ctgtgaagtt   | 720  |
| ctaccaagcc t  | caacaagtg  | aactttatgg   | gaaagtgcag   | gaaatacccc | agaaggagac   | 780  |
| cacccctttc t  | atccccggt  | caccctatgg   | ggcagcaaaa   | ctctatgcct | attggattgt   | 840  |
| ggtgaacttc c  | gtgaggcgt  | ataatctctt   | tgcagtgaac   | ggcattctct | tcaatcatga   | 900  |
| gagtcccaga a  | gaggagcta  | atttcgttac   | tcgaaaaatt   | agccggtcag | tagctaagat   | 960  |
| ttaccttgga c  | aactggaat  | gtttcagttt   | gggaaatctg   | gatgccaaac | gagattgggg   | 1020 |
| ccatgccaag g  | actatgtgg  | aggctatgtg   | gttgatgttg   | cagaatgatg | agccggagga   | 1080 |
| cttcgttata g  | ctactgggg  | aggtccatag   | tgtccgggaa   | tttgtcgaga | aatcattctt   | 1140 |
| gcacattgga a  | aaaccattg  | tgtgggaagg   | aaagaatgaa   | aatgaagtgg | gcagatgtaa   | 1200 |
| agagaccggc a  | aagttcacg  | tgactgtgga   | tctcaagtac   | taccggccaa | ctgaagtgga   | 1260 |
| gaccaattaa t  | tgttcaaag  | gtaatatata   | gagatctcaa   | gagcatttac | tttggcacag   | 1320 |
| tetetgeece t  | ttgtggtct  | ggctaacaat   | aaagctaaga   | tttgttttgt | aatgacggtt   | 1380 |
| ccagttggac t  | ggaatteet  | ttgaagttcc   | tgaagtttgc   | tgttgtgaga | gctcaagata   | 1440 |
| ttatttgact g  | gcaatgaaac | actctctgtg   | tattatcaaa   | atatogtata | cactgaactg   | 1500 |
| taacaacttg o  | agtaatcag  | acttctcctt   | cgaggtggcc   | tgcgtgagta | gagagcatcc   | 1560 |
| acagagagat g  | gagettetee | caacatccat   | ttagtgttca   | gtcatcacgg | gccactcatg   | 1620 |
| gatctgaacc t  | tcttacaag  | gtcgatacca   | acaggtatga   | ggagcagaga | gcagatcagc   | 1680 |
| acggagcagc a  | attagctggg | gcctccttgc   | ctggcctggc   | agtgaagata | gagcaccatg   | 1740 |
| ccagttgttc a  | agcttctcag | cgcttgcatc   | agetgaeeta   | aattatttaa | ggatgttgtc   | 1800 |
| acctctaaat a  | aaatatagtc | attgtcctgg   | tctctattta   | tattttaaaa | ctctttccca   | 1860 |
| gcccaccccg    | atggcctaaa | acaattcata   | tacttcagta   | gaagagggtg | tcacagggct   | 1920 |
| atttgcacag g  | gtttggatcc | agtgccagcg   | ; ttagccttca | cacaagttac | : actcctctga | 1980 |
| gatgtaactg    | tctcctgtgt | atgagtgcca   | a aagtatttga | taaccgccat | ctgtctgctc   | 2040 |
| tcaacacagg    | tttcacaago | : acagtgatgt | cttccattgt   | gtgcgtaaca | actgtgtgtg   | 2100 |
| tgcggagaga    | gaggactctg | tacgcatgca   | a catgtttgga | gctacataaa | a tetgagaage | 2160 |
| agttttgtct    | agtgtcgttt | cagtaaatg    | g gactcattta | gtcagtttg  | tttgctgtca   | 2220 |

| gaatcatatt                                      | ccagatatca | gaatgtgcta | tggcaggtac  | ttgtttaaaa | caagtaataa | 2280 |
|---|------------|------------|-------------|------------|------------|------|
| tatccttgta                                      | atgtttagca | atacagetga | ctggtctgtg  | ttccattact | tgtgtaaaag | 2340 |
| gaaatatgta                                      | aagcttttaa | aaataagttt | aaatgatcaa  | aagagatgcc | tgaaaatata | 2400 |
| ttttggcaca                                      | ggaaattaca | ggaatttata | aatgtgtcat  | gtggaaagat | aattacaaag | 2460 |
| tttattttaa                                      | tcttttatct | ctgactgaat | cttaagttaa  | aatattcttc | ccggcatcct | 2520 |
| gtctggtttt                                      | tgccactgtc | cctgctttac | gtgttacctt  | tccatgttga | tctgagttgc | 2580 |
| atttagttga                                      | ggaga      |            |             |            |            | 2595 |
| <210> 82<br><211> 170<br><212> DNA<br><213> Hom |            |            |             |            |            |      |
| _   | gacgctgcgc | cagtcaggcc | gcagccccgc  | tgcttggccc | gtcgggcccg | 60   |
| ccttggccgg                                      | ctcggcccgc | ccccggccct | ccctgcacgg  | cctcccgtgc | gcccctgtca | 120  |
| gactgtggcg                                      | gccggtcgcg | cggtgcgctc | tecetecetg  | cccgcagcct | ggagaggcgc | 180  |
| ttegtgetge                                      | acacccccgc | gttcctgccg | gcaccgcgcc  | tgccctctgc | cgcgctccgc | 240  |
| cctgccgccg                                      | accgcacgcc | cgccgcggga | catggcacac  | gcaccggcac | gctgccccag | 300  |
| cgcccggggc                                      | tccggggacg | gcgagatggg | caagcccagg  | aacgtggcgc | tcatcacegg | 360  |
| tatcacaggo                                      | caggatggtt | cctacctggc | tgagttcctg  | ctggagaaag | gctatgaggt | 420  |
| ccatggaatt                                      | gtacggcggt | ccagttcatt | taatacgggt  | cgaattgagc | atctgtataa | 480  |
| gaatccccag                                      | gctcacattg | aaggaaacat | gaagttgcac  | tatggcgatc | tcactgacag | 540  |
| tacctgcctt                                      | gtgaagatca | ttaatgaagt | aaagcccaca  | gagatctaca | accttggagc | 600  |
| ccagagccac                                      | gtcaaaattt | cctttgacct | cgctgagtac  | actgcggacg | ttgacggagt | 660  |
| tggcactcta                                      | cgacttctag | atgcagttaa | gacttgtggc  | cttatcaact | ctgtgaagtt | 720  |
| ctaccaagcc                                      | tcaacaagtg | aactttatgg | gaaagtgcag  | gaaatacccc | agaaggagac | 780  |
| cacccctttc                                      | tatccccggt | caccctatgg | ggcagcaaaa  | ctctatgcct | attggattgt | 840  |
| ggtgaacttc                                      | cgtgaggcgt | ataatctctt | tgcagtgaac  | ggcattctct | tcaatcatga | 900  |
| gagteccaga                                      | agaggagcta | atttcgttac | tcgaaaaatt  | agccggtcag | tagctaagat | 960  |
| ttaccttgga                                      | caactggaat | gtttcagttt | gggaaatctg  | gatgccaaac | gagattgggg | 1020 |
| ccatgccaag                                      | gactatgtgg | aggctatgtg | gttgatgttg. | cagaatgatg | agccggagga | 1080 |
| cttcgttata                                      | gctactgggg | aggtccatag | tgtccgggaa  | tttgtcgaga | aatcattctt | 1140 |
| gcacattgga                                      | aaaaccattg | tgtgggaagg | aaagaatgaa  | aatgaagtgg | gcagatgtaa | 1200 |

|   |  |            |            | 120        |              |            |      |
|---|--|------------|------------|------------|--------------|------------|------|
| ć | agagaccggc                               | aaagttcacg | tgactgtgga | tctcaagtac | taccggccaa   | ctgaagtggt | 1260 |
| ć | aaggactctc                               | tggccaccca | gtgcgtggcc | acgtttggct | ggctggctgg   | gcaaatgcgc | 1320 |
| į | catggaatg                                | cctggtgcca | gcctttggag | ctgtcagttc | agctccctgg   | catccttcag | 1380 |
| ( | gtgcatttc                                | caaaatcaga | acacagtgaa | ttcaatttga | gattatcatt   | cttaccacaa | 1440 |
| ě | acttcattgc                               | tctataaagg | aaaacaagag | ttaaacccac | caaaagaaat   | cattgatttc | 1500 |
| į | agaaatggaa                               | tctcacatca | cacgcagatc | tcctcatgtt | ggtctcctat   | tctaatggct | 1560 |
| 9 | ggtccctgat                               | aacctacagg | atgcctgact | taacaacgct | gtcccagacc   | tgcaggacat | 1620 |
|   | ctaggtcctc                               | tccagctatg | tctatgtctg | tgtctgtgta | ctcatacaca   | gtgcatactt | 1680 |
| • | gcacactata                               | ccataaatgg | tctccc     |            |              |            | 1706 |
|   | <210> 83 <211> 2667 <212> DNA <213> Homo | o sapien   |            |            |              |            |      |
|   |  | ttttttt    | gcttttttgt | tttttgagac | agggtcttgc   | tetgttgece | 60   |
|   | aggctggggt                               | gcagtggcga | gatctcagtt | cactgcaacc | tccacatccc   | ggctttaacc | 120  |
|   | gattctccca                               | tctcagcctc | acgagtagct | gggattacag | ggctgcatca   | acacacctgg | 180  |
|   | ctaatttttg                               | tatttttagt | acagatgggg | tttcaccatg | ttggccaggc   | tggtctcaaa | 240  |
|   | ctcctgacct                               | caggtgatct | gccagcttca | gcctcccaaa | gtgctgggat   | tacaggcgtg | 300  |
|   | agctaccacg                               | cctggccttg | gccatgtaga | tttctttcac | accttcctta   | ttccttcttc | 360  |
|   | agaagctgag                               | gccccatgag | gggaagctgg | tagccagggg | tcttttcagg   | tacctgtggc | 420  |
|   | caaattaggc                               | ccagatgtta | atgctctcat | ccatggctga | aaaaggccaa   | gaaactaaga | 480  |
|   | agagctgtgg                               | acgcacagac | tggcagagac | accgagaggg | cactgcgacc   | gcactgcaca | 540  |
|   | gatgagcaaa                               | ggaaggccaa | cttcagcatt | ccctcctcct | ctctaagccc   | ctccaaaagg | 600  |
|   | cctgctcagg                               | gagctctggc | caccgttctc | aagcaattgo | ctgacgtttt   | tactgacact | 660  |
|   | gagggctgga                               | gatggtccct | tttccaggca | gctatgtgtc | : catttcctag | agagggaaac | 720  |
|   | tgagtcccag                               | agaaggcggg | tgtggctgga | ggccatgggt | cagtcatagg   | cataaccatg | 780  |
|   | ccgtgaataa                               | ccgtcttggc | actctctgac | ttctttccct | ggcaatacaa   | cggatcccag | 840  |
|   | tgcccatttg                               | tcacataagg | aaactgaggc | tggcgtcttc | gatgaagcct   | ggcaggagtg | 900  |
|   | ggggtcacac                               | acggagtgca | aggctgtcca | agccttcaag | g tcattctga  | ctgatctttg | 960  |
|   | agccttagca                               | atacaggacg | tgggtccaag | agcatttcca | gcacaaaatg   | gcgcctcata | 102  |

gaaaaatgta gtcggagttc cagcgcccac ctccaccacg gctcaaagga gggggcgcgg

|            |              |              | 20.          |              |              |      |
|------------|--------------|--------------|--------------|--------------|--------------|------|
| geggeagege | acacttcatt   | tgcatataca   | aaaatcaacg   | gatttcgttg   | gctggcagcg   | 1140 |
| gtttgatctg | cacgaaactg   | ggtcgccctc   | tgaccggtct   | ctatcagtca   | ttggcccatg   | 1200 |
| cgagtctcgt | gacccaccag   | tcactaaagc   | gcatccctat   | tggcccgagc   | gccttccatc   | 1260 |
| acgcgtttaa | aaccccgagt   | cccgcctccc   | ctcgccacgc   | teeggeeeee   | aggaagaccg   | 1320 |
| ttagtcgcgt | gcgcgacttc   | ctcagcgtca   | gggcgcgccc   | gcgcgtatgc   | cgggccgtgg   | 1380 |
| cggcgtctca | ttcaaacggt   | ccaatcagcg   | gccgcgtcgc   | cttccccgtc   | ccccttcctc   | 1440 |
| tettettece | cttttcagcc   | cctccgttct   | ttgagaaccc   | tcccgacccc   | cacaaacccc   | 1500 |
| cattggccgg | cccgctaccc   | agccctctcc   | gccacttccc   | tcgcttctga   | ccatagtttg   | 1560 |
| cggggaaggg | agcgagcgcg   | tcgaaaacca   | aggaacgtgc   | gcgctgacgt   | cacggttgag   | 1620 |
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| cggggggaco | egececegge   | : cggccgcagc | : catgaactco | : aacgtggaga | acctacccc    | 1920 |
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| catcaaggto | tttcccaacg   | g aggaggacct | : caccgaccto | caggtcacca   | a togagggccc | 2040 |
| tgaggggaco | c ccatatgcts | g gaggtetgtt | ccgcatgaaa   | a creetgetg  | g ggaaggactt | 2100 |
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| cccgagcgg  | g egeegtgte  | a atgcccaaa  | c aaaaaacc   | t tctcgaaaa  | g gtgtcctcag | 2640 |
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<sup>&</sup>lt;210> 84 <211> 2648 <212> DNA <213> Homo sapien

<sup>&</sup>lt;400> 84

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| gattetecca teteageete acgagtaget gggattacag ggetgeatea acacacetgg | 180  |
| ctaatttttg tatttttagt acagatgggg tttcaccatg ttggccaggc tggtctcaaa | 240  |
| ctcctgacct caggtgatct gccagcttca gcctcccaaa gtgctgggat tacaggcgtg | 300  |
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| caaattaggc ccagatgtta atgctctcat ccatggctga aaaaggccaa gaaactaaga | 480  |
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| cotgotcagg gagototggo cacogitoto aagcaattgo otgaogitti tacigacaci | 660  |
| gagggctgga gatggtccct titccaggca gctatgtgtc catttcctag agagggaaac | 720  |
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| agecttagea atacaggaeg tgggtecaag ageattteca geacaaaatg gegeeteata | 1020 |
| gaaaaatgta gteggagtte cagegeeeae etecaceaeg geteaaagga gggggegegg | 1080 |
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| cgagtetegt gacceaceag teactaaage geatecetat tggeeegage geettecate | 1260 |
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| eggegtetea tteaaaeggt ceaateageg geegegtege etteceegte eccetteete | 1440 |
| tottottocc offiteagoo corcegitof tigagaacco tecegaceee egegggeeee | 1500 |
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| geggeggaeg teggaggeag eggggagegg ageggggeeg eeggggeete teeagggeeg | 1800 |
| cageggeage agttgggeee ceegeceegg eeggeggaee gaagaaeeta eeceegeaea | 1860 |
|   |      |

| tcatccgcct ggtgtacaag ga   | ggtgacga cactgaccgc | agacccaccc | gatggcatca | 1920 |
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| aggtctttcc caacgaggag ga   | cctcaccg acctccaggt | caccatcgag | ggccctgagg | 1980 |
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| gattetecca teteageete ac   | gagtagct gggattacag | ggctgcatca | acacacctgg | 180  |
| ctaatttttg tatttttagt ac   | agatgggg tttcaccatg | ttggccaggc | tggtctcaaa | 240  |
| ctcctgacct caggtgatct go   | cagettea geeteecaaa | gtgctgggat | tacaggcgtg | 300  |
| agctaccacg cetggeettg ge   | catgtaga tttctttcac | accttcctta | ttccttcttc | 360  |
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| caaattaggc ccagatgtta at   | geteteat ecatggetga | aaaaggccaa | gaaactaaga | 480  |
| agagetgtgg acgeaeagae tg   | gcagagac accgagaggg | cactgcgacc | gcactgcaca | 540  |
| gatgagcaaa ggaaggccaa ct   | tcagcatt ccctcctcct | ctctaagccc | ctccaaaagg | 600  |
| cctgctcagg gagctctggc ca   | cogttoto aagcaattgo | ctgacgtttt | tactgacact | 660  |
| gagggetgga gatagtaget  |                     |            |            |      |
| gagggeegga gaeggeeeet ee   | tccaggca gctatgtgtc | catttcctag | agagggaaac | 720  |

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| ccgtgaataa | ccgtcttggc | acțetetgae   | ttctttccct  | ggcaatacaa   | cggatcccag   | 840  |
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| cgagcgcgat                                       | aagaagctgg | cggccaagaa | aaagacggac | aagaagcggg | cgctgcggcg | 2640 |
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| caactctgtc                                       | tctaagttat | ttaaattatg | gctggggtcg | gggagggtac | agggggcact | 2760 |
| gggacctgga                                       | tttgtttttc | taaataaagt | tggaaaagca |            |            | 2800 |
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| ccagaagcgc                                       |            |            |            |            |            | 120  |
|  |            |            |            | ctctgagggc |            | 180  |
| ggttaaagct                                       |            |            |            |            |            | 240  |
|  |            |            |            | caagtcttat |            | 300  |
|  |            |            |            |            |            | 360  |
|  |            |            |            | agcaccctgg |            |      |
| _  |            |            |            | acaaaatctt |            | 420  |
|  |            |            |            | accacttcaa |            | 480  |
| ggaggtgcta                                       | tcgcaccact | gtgtatttaa | ctgccttgtg | tacagttatt | tatgcctctg | 540  |
| tatttaaaaa                                       | actaacaccc | agtctgttcc | ccatggccac | ttgggtcttc | cctgtatgat | 600  |
| tccttgatgg                                       | agatatttac | atgaattgca | ttttacttta | atcacactgt | atgcgtgtgt | 660  |
| gggtgttttg                                       | tagggaaagc | tcttctcaga | gtggggagct | ggtgggtgtc | acagcct    | 717  |
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|  |            |            |            | cataacgctt |            | 180  |
|  |            |            |            | tagaattcac |            | 240  |
|  |            |            |            | ctacggggca |            | 300  |
|  | _          |            |            | ctccccttcc |            | 360  |
|  |            |            |            | cgtcagaggc |            | 420  |
|  | •          |            |            |            |            | 480  |
| rgagecacea                                       | gaaagtgaag | agetgeeaga | taatgttatg | gtttccaagc | cagegeeeta | 400  |

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                                                                       300
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                                                                        480
                                                                        540
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| ggttttaaaa                                       | aatacatatc | tggttttggt | aaggtatttt | taatcaatta | ggcttgtagt | 600 |
|--|------------|------------|------------|------------|------------|-----|
| atcagtgaaa                                       | tactgtaggt | ttagggactg | ggctagcttc | atatcagatt | tacttgttaa | 660 |
| gtgactgttt                                       | tggaatgttt | acttttggac | tgggtttgta | acacggttaa | aggcaatgag | 720 |
| aaacaagcag                                       | aattccagga | gtccttgaag | cagagggcac | tggaagacaa | tatagca    | 777 |
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| gtttgtaaca                                       | cggttaaagg | caatgagaaa | caagcagaat | tccaggagtc | cttgaagcag | 540 |
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| ttgtataccg                                       | cagattcagg | catggattcc | gtgaaggaac | aacacctaaa | cccaagaggg | 240 |
| cagctgttgc                                       | agcatccagt | tcatcttaag | aatgtcaacg | attagtcatg | caataaatgt | 300 |
| tctggtttta                                       | aaaaatacat | atctggtttt | ggtaaggtat | ttttaatcaa | ttaggcttgt | 360 |
| agtatcagtg                                       | aaatactgta | ggtttaggga | ctgggctagc | ttcatatcag | atttacttgt | 420 |
| taagtgactg                                       | ttttggaatg | tttacttttg | gactgggttt | gtaacacggt | taaaggcaat | 480 |
| gagaaacaag                                       | cagaattcca | ggagtccttg | aagcagaggg | cactggaaga | caatatagca | 540 |

| <210> 92<br><211> 1103<br><212> DNA              | 3          |            |            |            |            |      |
|--|------------|------------|------------|------------|------------|------|
|  | o sapien   |            |            |            |            |      |
| <400> 92<br>gatgcatgct                           | cgagcggcgc | aggtgatgga | tgcgcccggg | caggtggccg | agcaggaggc | 60   |
| gccatcatgg                                       | gagtggacat | ccgccataac | aaggaccgaa | aggttcggcg | caaggagccc | 120  |
| aagagccagg                                       | atatctacct | gaggctgttg | gtcaagttat | acaggtttct | ggccagaaga | 180  |
| accaactcca                                       | cattcaacca | ggttgtgttg | aagaggttgt | ttatgagtcg | caccaaccgg | 240  |
| ccgcctctgt                                       | ccctttcccg | gatgatccgg | aagatgaagc | ttcctggccg | ggaaaacaag | 300  |
| acggccgtgg                                       | ttgtggggac | cataactgat | gatgtgcggg | ttcaggaggt | acccaaactg | 360  |
| aaggtatgtg                                       | cactgcgcgt | gaccagccgg | gcccgcagcc | gcatcctcag | ggcagggggc | 420  |
| aagatcctca                                       | ctttcgacca | gctggccctg | gactccccta | agggctgtgg | cactgtcctg | 480  |
| ctctccggtc                                       | ctcgcaaggg | ccgagaggtg | taccggcatt | tcggcaaggc | cccaggaacc | 540  |
| ccgcacagcc                                       | acaccaagtg | agtatcaggc | ccccagccct | gccctctccc | cagactcagc | 600  |
| ctgcagggcc                                       | aggeetggee | acacttgggc | tgcttctcct | atccctcgta | ccagccagcc | 660  |
| ccaggccttc                                       | tggaactctc | agctgcctca | tcctgcctgc | ttttcccttc | aacactgctt | 720  |
| ggagctgtgt                                       | tttccagggc | ctgagcaggc | atctcctggg | gagcatgcag | ttcaccggcc | 780  |
| tttgccagcc                                       | caggctggga | ccctccaggt | ggtggggcag | gtgcttccat | agcccctcct | 840  |
| ggctccttgg                                       | cttcccactc | tgccaagctt | ttccagcagc | attgactctc | ctggggctga | 900  |
| atgtgacagg                                       | cttgtggtgt | tcctgtgcca | ctccacaatg | gcctccttta | agagggcccc | 960  |
| cctctcactc                                       | cctcctctcc | ccacagaccc | tacgtccgct | ccaagggccg | gaagttcgag | 1020 |
| cgtgccagag                                       | gccgacgggc | cagccgaggc | tacaaaaact | aaccctggat | cctactctct | 1080 |
| tattaaaaag                                       | atttttgctg | aca        |            |            |            | 1103 |
| <210> 93<br><211> 398<br><212> DNA<br><213> Homo | sapien     |            |            |            |            |      |
| <400> 93   |            |            |            |            |            |      |
|  |            |            | tgcgcccggg |            |            | 60   |
|  | •          |            | aaggaccgaa |            |            | 120  |
|  |            |            | gtcaagttat |            |            | 180  |
|  |            |            | aagaggttgt |            |            | 240  |
| guuluuu  |            | uatgatccoo | aagatgaagc | EECCEGGCCG | ggaaaacaag | 300  |

| acggccgcgg  | ccaragagac    | Cataactgat | gargracaga | cccaggaggc | acccaaaccg | 250  |
|---|---------------|------------|------------|------------|------------|------|
| atcggccgcg  | accacgctaa    | gccggattcc | agcacact   |            |            | 398  |
| <210> 94<br><211> 1673<br><212> DNA<br><213> Homo | 3<br>o sapien |            |            | ·          |            |      |
| <400> 94<br>gccagtgtga                            | tggattggtc    | gcggcgaggt | ggccgagcag | gaggcgccat | catgggagtg | 60   |
| gacatccgcc  | ataacaagga    | ccgaaaggtt | cggcgcaagg | agcccaagag | ccaggatatc | 120  |
| tacctgaggc  | tgttggtcaa    | gttatacagg | tttctggcca | gaagaaccaa | ctccacattc | 180  |
| aaccaggttg  | tgttgaagag    | gttgtttatg | agtcgcacca | accggccgcc | tctgtccctt | 240  |
| tcccggatgg  | tgagtggctg    | gtccagagag | cacggtagac | ctgggagccg | ctgggtcttg | 300  |
| tctgtctgga  | agggaggcag    | gacttggage | tctgggagca | accagggcat | caagggtttg | 360  |
| tctcagccag  | tcgcgagcgt    | agagcttgga | ctgttggttg | gaactgagtg | tccatgggca | 420  |
| gtgggcaagt  | ctcctggtcc    | tcctctcctt | ctctttgtga | ggtggaggtg | teetggtgge | 480  |
| ttcaggagat  | tgccccaagt    | cataactgag | ttttatgtca | aaggototgo | tgagggaggc | 540  |
| cctattgaac  | agtctgcatt    | tttttttta  | tctggagctt | tecceagetg | gacctgagga | 500  |
| gaggcctgtg  | gggtttaagg    | ggtgtggtgt | gagtggggg  | tcccgttaga | gggcacagct | 660  |
| gccctggcct  | ggggatgcct    | caagcggggc | tgaatgtaaa | caccagaaca | acttacgacg | 720  |
| tacatcctcc  | ccaccctaag    | atccggaaga | tgaagcttcc | tggccgggaa | aacaagacgg | 780  |
| ccgtggttgt  | ggggaccata    | actgatgatg | tgcgggttca | ggaggtaccc | aaactgaagg | 840  |
| tgagctggcg  | ggggctgggc    | agacccatca | gacccttgct | gtactgtgct | gtgctgtgct | 900  |
| gtgctgtgct  | gtgctgtgct    | ggtccatcct | cagtcttgcc | accgctggtg | aggcaagcat | 960  |
| cctgcctgtc  | tgagtggctt    | tgaggattgg | tgtgggtcgt | ctctggcatg | gttgatggtg | 1020 |
| gggtcccctg  | ggcaagcttc    | ttctggggca | agttgtaaac | tgttagtgct | ggcccctaga | 1080 |
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| gcccagcttc  | cttgctgtag    | tttctagcat | aacaccaggg | atgtggtggt | gaaagttccg | 1200 |
| gaagtgttga  | gaactgctgc    | tgtcagccca | gggtgeteeg | ggtagtaaac | gatagtgttg | 1260 |
| gttaccaggg  | ctctgctgcg    | ccaaggcagt | ggagggtgag | tcctggcagc | ccctgggtga | 1320 |
| ccacgcaggg  | gtctgacccg    | ccagccttgt | cctctccacc | aggtatgtgc | actgcgcgtg | 1380 |
| accagccggg  | cccgcagccg    | catcctcagg | gcagggggca | agateeteae | tttcgaccag | 1440 |
| ctggccctgg  | actcccctaa    | gggctgtggc | actgtcctgc | tctccggtcc | tcgcaagggc | 1500 |

| cgagaggtgt                             | accggcattt | cggcaaggcc | ccaggaaccc | cgcacagcca | caccaaaccc | 1560 |
|--|------------|------------|------------|------------|------------|------|
| tacgtccgct                             | ccaagggccg | gaagttcgag | cgtgccagag | gccgacgggc | cagccgaggc | 1620 |
| tacaaaaact                             | aaccctggat | cctactctct | tattaaaaag | atttttgctg | aca        | 1673 |
| <210> 95 <211> 112 <212> DNA <213> Hom |            |            |            |            |            |      |
| <400> 95<br>tgagtgcctg                 | gaccctgacc | tccaacggga | gatecttece | ggggccattc | cctaaatcgt | 60   |
| gggcttgttt                             | cttgctccct | ctggctattc | tgtgtccctg | tggctgctcc | ccaaccctaa | 120  |
| gggctgtccc                             | tgatctctct | tgctatttcc | ccaagcccga | gactgctccc | ttacaaattc | 180  |
| tggctgctcc                             | ctttccctgt | gtccaatacc | gtgtgatcac | ctgtccctcc | ctggtccctc | 240  |
| taattctccc                             | ttgtgactat | tctgtaatcc | cactccccgt | tectgagece | ccgggattgt | 300  |
| ttcttggttc                             | cccagaatgt | tctccaagaa | cccaaagtgc | tgttccctaa | tcccactgac | 360  |
| ccttcctcag                             | cacatctgct | tgttgctctg | cagctctgac | tattccagac | accagagggt | 420  |
| gttacactga                             | cctcctggct | agtccaagcc | tgttccctga | tggctcctga | tggcctctga | 480  |
| gcccagtgac                             | tatcttgaag | ccccactggc | tattcctgtg | cttaaactcc | ttagccgcca | 540  |
| accgggtggg                             | gccccttact | actgccagtt | tttccctgtc | ctttgggaca | attccacagg | 600  |
| cgacctgcct                             | gactgacccc | cagggcctat | tccccagcca | ccattccagg | acactgattt | 660  |
| actggcttac                             | tgttaccttt | caggacccag | tgacatttcc | ctcaatggga | ggggctgtcc | 720  |
| tggggtatgt                             | acttaggtgg | ggtgagagca | agtctcccca | geggeteagt | gttgaaagct | 780  |
| gaggagagga                             | agttctgctc | cacgggggag | gcagggagaa | gccctcagag | atgtttccct | 840  |
| cctcccttcc                             | aggtgtttcc | ctgtttatcc | atcccctgca | aactgcagag | tggcactcat | 900  |
| tgcttgtgga                             | cggaccagct | cctccaaggc | tctgaaaagg | gcttccagtc | ccgtcacctt | 960  |
| gcctgcctgc                             | ctcgggagcc | agggctgtgc | acctggcagt | ccctgcggtc | ccagatagcc | 1020 |
| tgaatcctgc                             | ccggagtgga | agctgaagcc | tgcacagtgt | ccaccctgtt | cccactccca | 1080 |
| tctttcttcc                             | ggacaatgaa | ataaagagtt | accacccagc | a          |            | 1121 |
| <210> 96 <211> 570 <212> DNA <213> Hom | •          |            |            |            |            |      |
| <400> 96 ccagtccctg                    | gggcgcggcc | tagcaaccac | gagggggcga | ggctctgatg | ggaatggtcc | 60   |

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|---|--------------|------------|--------------|------------|------------|------|
| cactggaaat                                      | ggggaccacc   | ccaatttcag | r tctatcagaa | ggccgggcct | ttggcagctt | 120  |
| ggcagctcag                                      | J ccaatcacaa | gctgcttgtc | ggtccccgcc   | ccgcctttct | ccttaggaaa | 180  |
| actgcaggat                                      | ggactcttgc   | acatcactac | ctgcagtttt   | gtggctccct | ggaacagcct | 240  |
| gagcttagct                                      | cagcgccggg   | gcttcaccaa | gacctacact   | gttggctgtg | aggaatgcac | 300  |
| agtgtttccc                                      | : tgtttatcca | tcccctgcaa | actgcagagt   | ggcactcatt | gcttgtggac | 360  |
| ggaccagcto                                      | ctccaaggct   | ctgaaaaggg | cttccagtcc   | cgtcaccttg | cctgcctgcc | 420  |
| tcgggagcca                                      | gggctgtgca   | cctggcagtc | cctgcggtcc   | cagatagcct | gaatcctgcc | 480  |
| cggagtggaa                                      | gctgaagcct   | gcacagtgtc | caccctgttc   | ccactcccat | ctttcttccg | 540  |
| gacaatgaaa                                      | taaagagtta   | ccacccagca |              |            |            | 570  |
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| gatacagagg                                      | cageggeegg   | cgccccggga | tetgggggcg   | caccctcagg | gtcgcagggg | 120  |
| gtgctgatcg                                      | gggacaggct   | gtactccggg | gtgctcatca   | ccttggagaa | ctgcctcctg | 180  |
| cctgacgaca                                      | agctccgttt   | cacgccgtcc | atgtcgagcg   | gcctcgacac | cgacacagag | 240  |
| accgacctcc                                      | gcgtggtggg   | ctgcgagctc | atccaggcgg   | ccggtatcct | gctccgcctg | 300  |
| ccgcaggtgg                                      | ccatggctac   | cgggcaggtg | ttgttccagc   | ggttctttta | taccaagtcc | 360  |
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| gaagaggccc                                      | caagacgcat   | acgggacgtc | atcaatgtgt   | ttcaccgcct | tcgacagetg | 480  |
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| caaattataa                                      | aggcggaaag   | acgagttctc | aaagagttgg   | gtttctgcgt | ccatgtgaag | 600  |
| catcctcata                                      | agataatcgt   | tatgtacctt | caggtgttag   | agtgtgagcg | taaccaacac | 660  |
| ctggtccaga                                      | cctcatgggt   | agcctctgag | ggtaagtgac   | taagacttct | cctctgctgt | 720  |
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| ccactgtctg | tgaagaacac | caagaggagg | ctggagggcg | ccaagaaagc | caaggcggac | 3060 |
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| cttcatga                      | acc ggattcatcc                    | tgttactggt   | atteggetet | gtgacagtct | accatattat | 240 |
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| cactccac                      | ctg tgcaatgctg                    | gccctgcacg   | ctggggctgt | tgcccctgcc | cccttggtcc | 360 |
| tgccccta                      | aga tacagcagtt                    | tatacccaca   | cacctgtcta | cagtgtcatt | caataaagtg | 420 |
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| gtcaagct                      | tcc cagttgctct                    | tgatcctggg   | gccaagattt | cagtcattgt | ggaaacagtc | 180 |
| tacaccca                      | atg tgcttcatcc                    | atatccaacc   | cagatcaccc | agtcagagaa | acagtttgtg | 240 |
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|                               | gca ctgctcttcc                    |              |            |            |            | 120 |
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180

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| aaaatcagtg              | ccctccctgt    | tgctctagga | ggctcctgct | ggcttggtag | aagacagaat | 900  |
| tcgatctgcc              | tgtccctttt    | teceetgggg | tttgacacac | aggeteetet | cagcatgagg | 960  |
| tggagcagtg              | accaggtgga    | gcagtgacca | ggacgcctct | ggcccagtgc | tgcccagcct | 1020 |
| ccccgcccgc              | teccaggege    | cccatgtcct | cacaggccag | gacgccatgg | caggatggag | 1080 |
| aggacttggt              | ggatttttgt    | ttettgeetg | acctcagttt | catgaaagaa | agtggaagct | 1140 |
| acagaattat              | tttctaaaat    | aaaggctgaa | ttgtctgaaa | aa         |            | 1182 |
|                         | 5<br>o sapien |            |            |            |            |      |
| <400> 105<br>agcgggcgcg | ctccgctcgg    | cagcctgtgg | gacgcgaccg | cggcgctagt | ctcgttcctt | 60   |
| tgtgctgcgg              | cggcggcttc    | tcgagtcctc | cccgacgcgt | cctctaggcc | agcgagcccc | 120  |
| gcgctctccg              | gtgacggacc    | atgtcggcgg | cgggagcggg | cgcgggcgta | gaggcgggct | 130  |
| tctccagcga              | ggagctgctc    | tcgctccgtt | tcccgctgca | ccgcgcctgc | cgcgacgggg | 240  |
| acctggccac              | gctctgctcg    | ctgctgcagc | agacacccca | cgcccacctg | gcctctgagg | 300  |
| actccttcta              | tggctggacg    | cccgtgcact | gggccgcgca | tttcggcaag | ttggagtgct | 360  |
| tagtgcagtt              | ggtgagagcg    | ggagccacac | tcaacgtctc | caccacacgg | tacgcgcaga | 420  |
| cgccagccca              | cattgcagcc    | tttgggggac | atcctcagtg | cctggtctgg | ctgattcaag | 480  |
| caggagccaa              | cattaacaaa    | ccggattgtg | agggtgaaac | tcccattcac | aaggcagctc | 540  |

gctctgggag cctagaatgc atcagtgccc ttgtggcgaa tggggctcac gtcgacctga

600

| gaaatgccag tggcctgaca | gcagcagaca | ttgcacaaac | ccagggtttc | caagagtgtg | 660  |
|-----------------------|------------|------------|------------|------------|------|
| cccagtttct cttgaacctc | cagaattgtc | atctgaacca | tttctataac | aatggcatct | 720  |
| taaatggggg tcatcagaat | gtatttccta | atcatattag | tgtgggaaca | aatcgaaaga | 780  |
| gatgcttgga agactcagaa | gactttggag | taaagaaagc | tagaactgaa | gctcaaagct | 840  |
| tggattctgc cgtgccactc | acgaatggcg | acacagaaga | cgatgctgac | aaaatgcacg | 900  |
| ttgataggga gtttgctgtt | gtaacaggtg | ggagtggaca | gtttcctgtt | agctgcaaca | 960  |
| acaatccaat ggttgaagac | accaaacagc | aggagagtgg | ttctgttgga | ccaaaagaaa | 1020 |
| tagaaatata tactgtgtca | gcaatgcaga | cccctgtcg  | ttgcaggaat | caatatgaaa | 1080 |
| aacagtagct ccgtatcgaa | tacattgaca | aatggatgtg | tcatcaatgg | acatttggac | 1140 |
| ttcccctcca cgaccccgct | cagtgggatg | gaaagcagga | atggccagtg | cttgacagga | 1200 |
| actaacggaa ttagcagtgg | attagcccca | ggacagccgt | ttccgagtag | ccagggttct | 1260 |
| ctctgcatta gtgggactga | ggagccagag | aagaccctga | gagctaaccc | tgagttgtgc | 1320 |
| ggttctctgc acctgaacgg | gagtccaagt | agctgcatag | ccagtaggcc | ttectgggtg | 1380 |
| gaagacattg gggataacct | gtactatgga | cactaccacg | ggtttgggga | cactgctgaa | 1440 |
| agcatcccag aactgaacag | tgtggtcgag | cattccaagt | ccgtgaaggt | gcaggagcgg | 1500 |
| tacgacagtg ccgtgctggg | caccatgcac | ctgcaccacg | gctcctagag | acgctgacct | 1560 |
| ggctctcgga aacgcaggag | tecttectgg | tagccagctc | agaataccca | tgtagcagca | 1620 |
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| acttcgacaa gttcccagca | actgcttgtt | tgtgcatgag | tagggcttac | taagtgcata | 1740 |
| gatgtttcta cagtgaggtg | tcctttttat | aaggtgcact | tttggagttt | ttctgatgcc | 1800 |
| aatctcaaca ttgtcttttt | aatactgtca | ccagatattg | ccatttttct | ttttgttaaa | 1860 |
| agattatatg atcaagataa | attggggtgg | taaatcaggt | gcctggtaat | ttatctcttt | 1920 |
| gcacatgggc atcattttaa | aaagcttgct | tccactcttt | tctgtagaat | ttgacggaac | 1980 |
| acagetattt ceetatgeaa | ggtacagcct | tacaaagatt | tctgcagtga | tttgtgtgaa | 2040 |
| gaagagaatg tttgtctttt | tcaatgaagc | tttgcagatc | accatgtggt | tgaaggtttt | 2100 |
| agttgtggac acagtggtcc | ctccttaatg | atgaagatca | ctgccttggg | cttcatggaa | 2160 |
| aacaggccca gcctggggct | gcgtttggat | ttattgtttt | tattccacac | ttcctacttg | 2220 |
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| gtagaaactg tcaggtcaaa | atatttaact | gactgttgac | atgtattttc | ttttttcctt | 2340 |
| gtttttgttt tttggggttt | tctgctttaa | gatatatacc | actatgtata | tccagttaac | 2400 |
| tgagagaatt ttgactctct | taataaaact | gcattaagtt | tttgattttg | tagaaattag | 2460 |

| cttttgtcta   | ggcaactagt | ggttatactc | tgcaaatatt                              | gtaatgaatt   | ttactttt   | 2520 |
|--|------------|------------|---|--------------|------------|------|
| tgatttttgt   | aataaaaatt | ggtgcagata | aaatgtcaaa                              | tgaacaaacc   | agtgttctaa | 2580 |
| gagtgttact   | aacattttgt | tctaaaactg | tccttcacaa                              | attgaataaa   | aaactctcac | 2640 |
| actca  |            |            |   |              |            | 2645 |
| <210> 106<br><211> 2439<br><212> DNA<br><213> Homo | o sapien   |            |   |              |            |      |
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|  |            |            |   |              |            | 120  |
|  |            |            |   | ttttctcact   |            | 130  |
|  |            |            |   | gaaaagagct   |            |      |
|  |            |            |   | ttttttaatt   |            | 240  |
| atttatttga   | tacatgtatt | gtagttttag | ttgaattaaa                              | ctagttagta   | aaactgttgg | 300  |
| tgaggtagtg   | tttcagttac | tctgtagata | agctcttttg                              | ttaaaacctc   | ttaagttttt | 360  |
| tgttcctgaa   | tatgttacaa | acatctattt | tctcttactg                              | tagctatcac   | atgaggaaaa | 420  |
| aaacttgaaa   | aaaagacttc | atagtgtgtg | tttatgtttg                              | attttagaag   | acccttttga | 480  |
| ggacttcttt   | gggaatcgaa | ggggtccccg | aggaagcaga                              | agccgaggga   | cggggtcgtt | 540  |
| tttctctgcg   | ttcagtggat | ttccgtcttt | tggaagtgga                              | ttttcttctt   | ttgatacagg | 600  |
| atttacttca   | tttgggtcac | taggtcacgg | gggcctcact                              | tcattctctt   | ccacgtcatt | 660  |
| tggtggtagt   | ggcatgggca | acttcaaatc | gatatcaact                              | tcaactaaaa   | tggttaatgg | 720  |
| cagaaaaatc   | actacaaaga | gaattgtcga | gaacggtcaa                              | gaaagagtag   | aagttgaaga | 780  |
| agatggccag   | ttaaagtcct | taacaataaa | tggtgtggcc                              | gacgacgatg   | ccctcgctga | 840  |
| ggagcgcatg   | cggagaggcc | agaacgccct | gccagcccag                              | cctgccggcc   | teegeeegee | 900  |
| gaagccgccc   | cggcctgcct | cgctgctgag | acacgcgcct                              | cactgtctct   | ctgaggagga | 960  |
| gggcgagcag   | gaccgacctc | gggcacccgg | gccctgggac                              | cecetegegt   | ccgcagcagg | 1020 |
| attgaaagaa   | ggtggcaaga | ggaagaagca | gaagcagaga                              | gaggagtcga   | agaagaagaa | 1086 |
| gtcgaccaaa   | ggcaatcact | agaccggact | tgaggcacgc                              | ggtgcacccc   | cagacgctgg | 1140 |
| cgctccaccg   | tgctcggcat | geggtegtge | acacgcgcta                              | ggtagcagcg   | tcggtcagga | 120  |
| ctgtctcgag   | gccacactcg | ctcggcagga | ttatgcgatc                              | : acggatcagt | cagagcaggg | 126  |
| tcaggagacg   | gggctgacgg | cacgggtggc | ggggacagac                              | gtttgggact   | tggccgcgac | 132  |
| + a + a + a a + + a                                |            | tanntataat | ~~~ * * * * * * * * * * * * * * * * * * | ataatactta   | cagatectet | 138  |

| tcattcttt               | cggctactca    | accactccgc | atgctgctgg | aatatttctg | gctttagaag | 1440 |
|-------------------------|---------------|------------|------------|------------|------------|------|
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| cgtagtgttt              | ggaattagaa    | ggtaattcag | tagagtgtaa | cttagagaat | attgcaagtg | 1560 |
| acacattgaa              | tcctgcccgt    | cagggcacct | tttcctcaga | gcaatccggc | cacacgaata | 1620 |
| gaaggctgtc              | gtgaatcaca    | tcagatgtaa | aatcattcct | tctgtttact | cttttaattt | 1680 |
| tcatcctttg              | caggtagtgc    | aaattcaact | tcaaatatgg | tgtaggtttt | gctagattcc | 1740 |
| atatttttt               | cttggatttt    | tgctaattat | ttttagcaaa | aaatttttgc | tcagtggcac | 1800 |
| cctccctagt              | gtccatgggt    | tagggccatg | ctggggaaaa | cgggccggta | tttacacacg | 1860 |
| cgcaaaacac              | ccagagacgg    | cacaaggagg | ttgaactcat | gtttcagttc | gcgaacattg | 1920 |
| actccttacg              | aaagtcactt    | cattctaact | agatgegeee | acttccggtc | attatttcgt | 1980 |
| ttgcatgatg              | tattgcttct    | tcacgttttg | tttttattga | gcacggagta | gaattccagg | 2040 |
| gctgccttga              | cttcttccct    | gcatgctccc | tcccagtgac | tttccttccc | tttcacatga | 210  |
| ggatctgccg              | ttcatgttgc    | tttctccttt | gtcctcttgg | acttgagggc | attgtgaaaa | 216  |
| gctttgctgt              | gatttaaaaa    | tgccagcaat | tttaatctag | cagtgttgaa | gctgggaatt | 222  |
| ttttggcgca              | atccatgtag    | cagtgaccca | ggcttgggag | ccagaaacaa | gtgtgacctg | 228  |
| ggattttatt              | taacacaact    | gttgccaaag | agttggcttt | gtttatttgg | ttttggcggg | 234  |
| gagaggagtg              | gtatttgatg    | ctttctgtgg | acaatgtaac | cctaaacaca | tcatgtattt | 240  |
| taaatgccac              | ctacataaat    | aaaacataag | catattgaa  |            |            | 243  |
|                         | 3<br>o sapien |            |            |            |            |      |
| <400> 107<br>ggcagggcct | caggeceeca    | agagggctgt | gagcaggcac | cttcctgtga | ggcacgatgc | 6    |
| caccatgtga              | agttgtgtgg    | gtctttggag | agttctggca | ggttgagggc | ctgtgcgtgt | 12   |
| gggacccagg              | cctgccctgc    | ctgctgccgg | ccatgttccc | tctgcccagt | catgttgaca | 18   |
| attgtggcca              | aattcaacat    | cgggcctgtt | tcctgcaccc | acgttggcac | acactgagtt | 24   |
| caagacagtt              | tctgcagaaa    | tgaatccaga | ggtgtttatt | tctggcacgt | tggatcagac | 30   |
| gṛtcagaggg              | tgggcgtctt    | tctccatgca | cttcagctgg | atcagtcagt | gtcagccatt | 36   |
| gactttagca              | acagggaaat    | gaaagacttt | cctcaataag | aaatgggact | ttctcacgct | 42   |
| ggatggcaca              | gagctggcaa    | acatgaggcc | agtccacacc | aaggctttgc | ctgcagaaag | 48   |

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180

| ttgaaagaag gtggcaagag gaagaagcag aagcagagag aggagtcgaa gaagaagaag           | 600  |
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| tcgaccaaag gcaatcacta gaccggactt gaggcacgcg gtgcaccccc agacgctggc           | 660  |
| gctccaccgt gctcggcatg cggtcgtgca cacgcgctag gtagcagcgt cggtcaggac           | 720  |
| tgtctcgagg ccacactcgc tcggcaggat tatgcgatca cggatcagtc agagcagggt           | 780  |
| caggagacgg ggctgacggc acgggtggcg gggacagacg tttgggactt ggccgcgact           | 840  |
| ctctgcttct ctccagctct caatctgctg cattttcctc tagtgcttcc ggatcctctt           | 900  |
| cattetttte ggetaeteaa eeaeteegea tgetgetgga atatttetgg etttagaagt           | 960  |
| acaggagggc gcagatggct aactgagtaa cattcatgaa atgaggcttt ctgtggcggc           | 1020 |
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| cacattgaat cotgooogto agggoacott ttootcagag caatcoggoo acacgaatag           | 1140 |
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| catcetttge aggtagtgea aatteaaett caaatatggt gtaggttttg etagatteea           | 1260 |
| tatttttttc ttggattttt gctaattatt tttagcaaaa aatttttgct cagtggcacc           | 1320 |
| ctccctagtg tccatgggtt agggccatgc tggggaaaac gggccggtat ttacacacgc           | 1380 |
| gcaaaacacc cagagacggc acaaggaggt tgaactcatg tttcagttcg cgaacattga           | 1440 |
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| tgcatgatgt attgcttctt cacgttttgt ttttattgag cacggagtag aattccaggg           | 1560 |
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| tttggcgcaa tccatgtagc agtgacccag gcttgggagc cagaaacaag tgtgacctgg           | 1800 |
| gattttattt aacacaactg ttgccaaaga gttggctttg tttatttggt tttggcgggg           | 1860 |
| agaggagtgg tatttgatgc tttctgtgga caatgtaacc ctaaacacat catgtatttt           | 1920 |
| aaatgccacc tacataaata aaacataagc atattgaa                                   | 1958 |
| <210> 108<br><211> 3400<br><212> DNA<br><213> Homo sapien                   |      |
| <400> 108 ccgccccgtc gcgccagcgt gcggagaaac ctggcgccgc gaccagtgta tcgggggtct | 60   |
| tggaaatgge taagaaaget ggeetetgee taggaggete teggeagggg ggetgeeaat           | 120  |

ccgggatggt taccggaaat gaaccgcgcg atctggccct ttcacaccca cttagctttg

| tagggggttt | gtagtcttta | caggacaaaa | gggcacctat | tccagagcac | accccctttt   | 240  |
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| cttcctctcc | cttctgtgcc | tcactgcttt | ccgatctcat | agttgctccc | agtaagtgct   | 300  |
| ccaaagtcgt | gtggggccgc | atctgaggtc | tgcccgcctg | aattgggttg | gaggggttca   | 360  |
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| aggcgtgcct | ggtatcttca | tcgtaatagc | agagtcgtca | actttctatg | aaaggtggtt   | 480  |
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| gaagacttct | gaataaggtt | attctgtgtc | tttcatagta | gaaaccttaa | tgatcggtct   | 660  |
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| ggaatattat | agtattaaca | tctggaaaac | taggtaaatt | tatcttagaa | ttaagttttt   | 900  |
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| tataacaaac | teegecaace | attttttcag | aagaggtcag | aattgatcgc | caaaatccca   | 1080 |
| aatttttggg | taacaacatt | tgtcaaccat | ccacaagtgt | ctgcactgct | tggggaggaa   | 1140 |
| gatgaagagg | cactgcatta | tttgaccaga | gttgaagtga | cagaatttga | agatattaaa   | 1200 |
| tcaggttaca | gaatagattt | ttattttgat | gaaaatcctt | actttgaaaa | taaagttctc · | 1260 |
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| tacttggttc | ccgatatgga | tgatgaagaa | ggagaaggag | aagaagatga | tgatgatgat   | 1560 |
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| tgcagtcttt | tttttttt   | tttttttt   | ccctcttgtg | ctcagtcgcc | ctgttcttga   | 1800 |
| ggtctctttt | ctctactcca | tggttctcaa | tttatttggg | gggaaatacc | ttgagcagaa   | 1860 |
| tacaatggga | aaagagtctc | tacccctttc | tgttcgaagt | tcatttttat | cccttcctgt   | 1920 |
| ctgaacaaaa | actgtatgga | atcaacacca | ccgagctctg | tgggaaaaaa | gaaaaacctg   | 1980 |

| ctcccttcgc tctgctggaa | gctggagggt | gctaggcccc | tgtgtagtag | tgcatagaat | 2040 |
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| gatatttgac agacatcctt | gcagtttaag | atgacacttt | taaaataaat | tctctcctaa | 2340 |
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| gaattgacat tototattgt | aattttgttc | ctgtttattt | ttaaattttc | tttttgtttc | 2460 |
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| aattaattgt gcctgccacc | accatccaac | agacctggtg | ctctaatgcc | aagttataca | 2940 |
| cgggacagtt gctggcatgt | cttcattggc | tatataaaat | gtggccaaga | agataggete | 3000 |
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| tagatggata gtatgtaatt | tctgcacagg | tototgttta | gtaaatacat | cactgtatac | 3360 |
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<sup>&</sup>lt;210> 109 <211> 2775

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 109

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| agcaagaagc | gattgaacac   | attgatgaag   | tacaaaatga | aatagacaga   | cttaatgaac   | 360  |
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| ccagagttga | agtgacagaa   | tttgaagata   | ttaaatcagg | ttacagaata   | gatttttatt   | 600  |
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| aacgttcgag | tcaaacgcag   | aataaagcca   | gcaggaagag | gcagcatgag   | gaaccagaga   | 780  |
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| aagaaggaga | aggagaagaa   | gatgatgatg   | atgatgaaga | ggaggaagga   | ttagaagata   | 960  |
| ttgacgaaga | aggggatgag   | gatgaaggtg   | aagaagatga | agatgatgat   | gaagggagg    | 1020 |
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| ctcaatttat | ttggggggaa   | ataccttgag   | cagaatacaa | tgggaaaaga   | gtctctaccc   | 1260 |
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| ggatttgat  | cttttctctc   | : agcataggta | tgcttactat | gaccttccaa   | gtttgacttg   | 1980 |
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| tattccatca   | ataggcaaaa | gtgtaacaac | ctatctagat | ggatagtatg | taatttctgc | 2700 |
| acaggtetet   | gtttagtaaa | tacatcactg | tataccgatc | aggaatcttg | ctccaataaa | 2750 |
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| cccgccacgc   | cgctgtcgcc | cacgcgcctg | tegeggetge | aggagaagga | ggagctgcgc | 130  |
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| gaccggctcc   | tgctcaagat | ctcagagaag | gaggaggtga | ccacgcgcga | ggtgagtggc | 300  |
| atcaaggcgc   | tgtacgagtc | ggagctggcc | gatgcccgga | gagtcctgga | tgagacggct | 360  |
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| cggctctaca | agctggagct | ggagcagacc | taccaggcca | agctggacag | cgccaagctg | 900  |
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| atgcgcctgg | agtccctcag | ctaccagete | teeggeetee | agaagcaggc | cagtgccgct | 1020 |
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| gcagtggcca a                                       | aaagcagct   | ggagaaggag   | acgctgatgc | gtgtggacct   | ggagaaccgc   | 660  |
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| ctgggcacgg | gcacgggtgg | cageggtgge | ttccacctgg | cccagcaggc | ctcggcctcg | 1440 |
| ggtagcgtca | gcatcgagga | gatcgacctg | gagggcaagt | ttgtgcagct | caagaacaac | 1500 |
| tcggacaagg | atcagtctct | ggggaactgg | agaatcaaga | ggcaggtctt | ggaggggag  | 1560 |
| gagatcgcct | acaagttcac | gcccaagtac | atcctgcgcg | ccggccagat | ggtcacggtg | 1620 |
| tgggcagctg | gtgcgggggt | ggcccacagc | ccccctcga  | cgctggtgtg | gaagggccag | 1680 |
| agcagctggg | gcacgggcga | gagcttccgc | accgtcctgg | ttaacgcgga | tggcgaggaa | 1740 |
| gtggccatga | ggactgtgaa | gaagtcctcg | gtgatgcgtg | agaatgagaa | tggggaggaa | 1800 |
| gaggaggagg | aagccgagtt | tggcgaggag | gatcttttcc | accaacaggg | ggacccgagg | 1860 |
| accacctcaa | gaggctgcta | cgtgatgtga | acccacactc | ctcatccaca | cacctttctt | 1920 |
| tacccagagc | cactgaaaac | tatttttata | tcattggctt | tctttagttc | ttgatacatt | 1980 |
| tctagagaat | ttctaagcga | actgccagaa | cgtgtgggtg | ggtctccccc | agccctccct | 2040 |
| cctggcgggt | ctcctccagc | ctcacttcgc | tgccacttcg | ccgctgcccc | ggagactttt | 2100 |
| caatcccacc | ccactcctca | tctcaccatt | tggtcaaatt | ggaagcccag | ggccaggacc | 2160 |
| cggaggttta | gaagatgctt | gggcttggag | ggaggagggc | cggcgaggct | agcgagggga | 2220 |
| caggagacgg | ccctgctgcg | gacggagcgc | ggaaactgcg | taggaattca | gtggtggtgg | 2280 |
| gtttttttaa | ggctttctac | aaaaccaaat | tcagaatcca | ggcgtcgacc | tggtggggcc | 2340 |
| cggggccaag | cctgcattct | ggctgcccag | cttcggacag | cgggaactcc | tcaggcagcc | 2400 |
| acgcagcggg | tgtgggccag | catggggatg | gegtggeeee | cagggegggt | tttcactccg | 2460 |
| ctgcctgggc | ttccagattc | ccgttctggc | agcgcaccgg | ccgggtttct | cggaccgttg | 2520 |
| actttatttg | ggggagtttt | cccgcagttc | agttcctgac | tgtgcaaggc | caacagggca | 2580 |
| ggggagggga | agacctgggg | aaggaagaat | gaggacagtc | ccgtcgtaag | acctgtcaca | 2640 |
| acaataagca | gggaggggag | atgtggaggg | gacacatctg | gttgccttgg | aggcagaagc | 2700 |
| tgtgagtttc | agaacagctg | tctgcaggga | acgccaccat | gttgaccctc | tggaggagag | 2760 |
| cgctgtggag | cccctcccgt | gttccagctc | cgtctgccct | gtgcctatat | atacacatgc | 2820 |
| gtctatccat | actgtgcttt | tatctgtgat | tttctcgctg | aaaccatgtt | tctcagacag | 2880 |
| gccaaggcca | cctgactcct | atcacgacgc | acccaagccc | ctcagtccag | cttcccaatg | 2940 |
| cctggcaccc | cccttcggca | atagctcacc | gtttacaccc | tccctcatag | atacacagaa | 3000 |
| gttattttt  | taatggatat | ttatttttt  | acattggtca | gtacacaggt | caggagetea | 3060 |
| cgccagggcc | ttgaggacag | gctgaccctc | ctccccgggg | tggcgtgggg | ctggggcacc | 3120 |

| ctccgacggc | agageeteet | tcagaaagtg | cagctcaagt | cttaaagaca | ccaaaactga | 3180 |
|------------|------------|------------|------------|------------|------------|------|
| gccatgggca | cgcgccgtct | ccgggccatg | gcgttcactg | cagggcgggg | geggeacege | 3240 |
| tcccctgtga | ctgcatcccg | cctccctggg | gacctgcctg | tggcaggaag | gaatgggggg | 3300 |
| ccccagcccc | aggccgggaa | ggagccagcg | gccgacaaag | cagaaacacc | cgctgctcca | 3360 |
| cgtagcccct | gctcgctgtc | cttgctctca | gaagtcccgg | tcccatgtag | atagagggg  | 3420 |
| gcgcatctta | ccaaagcatt | tcctcctgga | ggctacgccg | ctgtgctccc | agtcaggcgg | 3480 |
| ctggtaggga | gctttgcctg | ccccggggat | accetetgee | agccgctgga | agtgggaatg | 3540 |
| ctggcgacag | actgtgtcct | ctttcccacc | ttcatagcag | gaatcacccg | gacccgactg | 3600 |
| gctgggcttc | gtgctagcga | gggttctggg | ggtgggtett | ggtgatcttg | tcctatgggg | 3660 |
| agtctgcagt | ggtctcagcc | acatcctatg | tattttggct | ctggaggagc | aaagctgtat | 3720 |
| cctggagtig | gtctgtgatt | tgccgacagc | cttgcaggct | gggctcaggg | acaaagtccc | 3780 |
| ccccaaaacc | cgcaggtcct | cattcctggg | gcgtcgccgc | ctgcagcctc | ttccaagccc | 3840 |
| tgcgtccagc | gagcgtcaca | gcacaacctg | caaaaacgga | getgggetge | agetgggget | 3900 |
| ggcatggact | ttcatttcag | agattcggtt | tttaagaaga | tgcatgccta | atgtgttctt | 3960 |
| tttttttcc  | aatgatttgt | aatatacatt | ttatgactgg | aaacttttt  | gtacaacact | 4020 |
| ccaataaaca | ttttgatttt | a          |            |            |            | 4041 |
|            |            |            |            |            |            |      |

<210> 113

<211> 165

<212> PRT

<213> Homo sapien

<400> 113

Gly Gly Arg Asn Arg Gly Arg Ala Gly Ser Gln Gly Gly Arg Gly Gly 1 5 10 15

Asn Arg Pro Ala Ile Ala Arg Gly Ala Ala Gly Gly Gly Arg Asn 35 40 45

Arg Pro Ala Pro Tyr Ser Arg Pro Lys Gln Leu Pro Asp Lys Trp Gln 50 55 60

His Asp Leu Phe Asp Ser Gly Phe Gly Gly Gly Ala Gly Val Glu Thr 65 70 75 80

Gly Gly Lys Leu Leu Val Ser Asn Leu Asp Phe Gly Val Ser Asp Ala

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163

85 90

95

Asp Ile Gln Glu Leu Phe Ala Glu Phe Gly Thr Leu Lys Lys Ala Ala 105

Val His Tyr Asp Arg Ser Gly Arg Ser Leu Gly Thr Ala Asp Val His 120

Phe Glu Arg Lys Ala Asp Ala Leu Lys Ala Met Lys Gln Tyr Asn Gly

Val Pro Leu Asp Ala Ser Tyr Ile Pro Pro Leu Leu Gln Leu Leu Pro 155

Glu Asp Ser Leu Leu

<210> 114 <211> 164 <212> PRT <213> Homo sapien

<220>

<221> MISC\_FEATURE

<222> (4)..(4)

<223> X=any amino acid

<400> 114

Gly Ala Gly Xaa Ala Pro Gly Arg Leu Gln Gly Gly Arg Gly Gly 5

Ala Gln Ala Ala Arg Val Asn Arg Gly Gly Pro Ile Arg Asn 20

Arg Pro Ala Ile Ala Arg Gly Ala Ala Gly Gly Gly Arg Asn Arg 35

Pro Ala Pro Tyr Ser Arg Pro Lys Gln Leu Pro Asp Lys Trp Gln His 55

Asp Leu Phe Asp Ser Gly Phe Gly Gly Gly Ala Gly Val Glu Thr Gly 70

Gly Lys Leu Leu Val Ser Asn Leu Asp Phe Gly Val Ser Asp Ala Asp 85

Ile Gln Glu Leu Phe Ala Glu Phe Gly Thr Leu Lys Lys Ala Ala Val

105 110 100

His Tyr Asp Arg Ser Gly Arg Ser Leu Gly Thr Ala Asp Val His Phe 120

Glu Arg Lys Ala Asp Ala Leu Lys Ala Met Lys Gln Tyr Asn Gly Val 135

Pro Leu Asp Ala Ser Tyr Ile Pro Pro Leu Leu Gln Leu Leu Pro Glu 150 155

Asp Ser Leu Leu

<210> 115 <211> 256 <212> PRT

<213> Homo sapien

<400> 115

Met Ala Leu Arg Val Thr Arg Asn Ser Lys Ile Asn Ala Glu Asn Lys 5

Ala Lys Ile Asn Met Ala Gly Ala Lys Arg Val Pro Thr Ala Pro Ala 25

Ala Thr Ser Lys Pro Gly Leu Arg Pro Arg Thr Ala Leu Gly Asp Ile

Gly Asn Lys Val Ser Glu Gln Leu Gln Ala Lys Met Pro Met Lys Lys 50 · 55

Glu Ala Lys Pro Ser Ala Thr Gly Lys Val Ile Asp Lys Lys Leu Pro 65

Lys Pro Leu Glu Lys Val Pro Met Leu Val Pro Val Pro Val Ser Glu 90

Pro Val Pro Glu Pro Glu Pro Glu Pro Glu Pro Val Lys Glu 100 105

Glu Lys Leu Ser Pro Glu Pro Ile Leu Val Asp Thr Ala Ser Pro Ser 120 125 115

Pro Met Glu Thr Ser Gly Cys Ala Pro Ala Glu Glu Asp Leu Cys Gln 135 130

Ala Phe Ser Asp Val Ile Leu Ala Val Asp Ala Glu Asp 145 150 155 160

Gly Ala Asp Pro Asn Leu Cys Ser Glu Tyr Val Lys Asp Ile Tyr Ala 165 170 175

Tyr Leu Arg Gln Leu Glu Glu Glu Gln Ala Val Arg Pro Lys Tyr Leu 180 185 190

Leu Gly Arg Glu Val Thr Gly Asn Met Arg Ala Ile Leu Ile Asp Trp 195 200 205

Leu Val Gln Val Gln Met Lys Ser Val Cys Ala Gly Pro Val Cys Gly 210 215 220

Pro Ile Asp Gly Pro Ala Lys Leu Gly Ala Gln Ile Ala Gly Gly Pro 225 230 235 240

Ala Val Trp Pro Leu Lys Gly Pro Arg Gly Arg Trp Gly Thr Leu Ala 245 250 255

<210> 116 ·

<211> 250

<212> PRT

<213> Homo sapien

<400> 116

Ala Gly Ser Ser Arg Arg Ala Ala Ala Glu Arg Leu Leu Val Ser Ala 1 5 10 15

Gly Cys Arg Ser Leu Ala Gly Arg Ala Ser Gly Val Leu Leu Pro 20 25 30

Ala Glu Leu Leu Pro Gly Glu Glu Glu Ala Met Ala Leu Arg Val Thr 35 40 45

Arg Asn Ser Lys Ile Asn Ala Glu Asn Lys Ala Lys Ile Asn Met Ala 50 55 60

Gly Ala Lys Arg Val Pro Thr Ala Pro Ala Ala Thr Ser Lys Pro Gly 70 75 80

Leu Arg Pro Arg Thr Ala Leu Gly Asp Ile Gly Asn Lys Val Ser Glu 85 90 95

Gln Leu Gln Ala Lys Met Pro Met Lys Lys Glu Ala Lys Pro Ser Ala

166

100 105 110

Thr Gly Lys Val Ile Asp Lys Lys Leu Pro Lys Pro Leu Glu Lys Val 115 120 125

Pro Met Leu Val Pro Val Pro Val Ser Glu Pro Val Pro Glu Pro Glu 130 135 140

Pro Ile Leu Val Asp Thr Ala Ser Pro Ser Pro Met Glu Thr Ser Gly 165 170 175

Cys Ala Pro Ala Glu Glu Asp Leu Cys Gln Ala Phe Ser Asp Val Ile 180 185 190

Leu Ala Val Asn Asp Val Asp Ala Glu Asp Gly Ala Asp Pro Asn Leu 195 200 205

Cys Ser Glu Tyr Val Lys Asp Ile Tyr Ala Tyr Leu Arg Gln Leu Glu 210 215 220

Glu Glu Gln Ala Val Arg Pro Lys Tyr Leu Leu Gly Arg Glu Val Thr 225 230 235 240

Gly Asn Met Arg Ala Ile Leu Ile Asp Trp 245 250

<210> 117

<211> 406

<212> PRT

<213> Homo sapien

<400> 117

Met Glu Ala Ala Ala Val Thr Val Thr Arg Ser Ala Thr Arg Arg Arg 1 5 10 15

Arg Arg Gln Leu Gln Gly Leu Ala Ala Pro Glu Ala Gly Thr Gln Glu 20 25 30

Glu Gln Glu Asp Gln Glu Pro Arg Pro Arg Arg Arg Pro Gly Arg 35 40 45

Ser Ile Lys Asp Glu Glu Glu Glu Thr Val Phe Arg Glu Val Val Ser 50 55 60 Phe Ser Pro Asp Pro Leu Pro Val Arg Tyr Tyr Asp Lys Asp Thr Thr 65 70 75 80

Lys Pro Ile Ser Phe Tyr Leu Ser Ser Leu Glu Glu Leu Leu Ala Trp 85 90 95

Lys Pro Arg Leu Glu Asp Gly Phe Asn Val Ala Leu Glu Pro Leu Ala 100 105 110

Cys Arg Gln Pro Pro Leu Ser Ser Gln Arg Pro Arg Thr Leu Leu Cys 115 120 125

His Asp Met Met Gly Gly Tyr Leu Asp Asp Arg Phe Ile Gln Gly Ser 130 135 140

Val Val Gln Thr Pro Tyr Ala Phe Tyr His Trp Gln Cys Ile Asp Val 145 150 155 160

Phe Val Tyr Phe Ser His His Thr Val Thr Ile Pro Pro Val Gly Trp 165 170 175

Thr Asn Thr Ala His Arg His Gly Val Cys Val Leu Gly Thr Phe Ile 180 185 190

Thr Glu Trp Asn Glu Gly Gly Arg Leu Cys Glu Ala Phe Leu Ala Gly 195  $\phantom{-}200\phantom{0}$  205

Asp Glu Arg Ser Tyr Gln Ala Val Ala Asp Arg Leu Val Gln Ile Thr 210 215 220

Gln Phe Phe Arg Phe Asp Gly Trp Leu Ile Asn Ile Glu Asn Ser Leu 225 230 235 240

Ser Leu Ala Ala Val Gly Asn Met Pro Pro Phe Leu Arg Tyr Leu Thr \$245\$ \$250\$

Thr Gln Leu His Arg Gln Val Pro Gly Gly Leu Val Leu Trp Tyr Asp 260 265 270

Ser Val Val Gln Ser Gly Gln Leu Lys Trp Gln Asp Glu Leu Asn Gln 275 280 285

His Asn Arg Val Phe Phe Asp Ser Cys Asp Gly Phe Phe Thr Asn Tyr 290 295 300

Asn Trp Arg Glu Glu His Leu Glu Arg Met Leu Gly Gln Ala Gly Glu 305 310 315 320

Arg Arg Ala Asp Val Tyr Val Gly Val Asp Val Phe Ala Arg Gly Asn 325 330 335

Val Val Gly Gly Arg Phe Asp Thr Asp Lys Val Gly Gly Phe Arg 340 345 350

Pro Arg Ala Ser Gly Pro Val Pro Pro Leu Gly Pro His Phe Leu Met 355 360 365

Asp Leu Pro Phe Pro Ser Ala Pro Gln Arg Asn Asp Ser Ser Cys Ser 370 375 380

Ser Gln Ser Gly Asp Pro Val Ala Leu Arg Asn Arg Cys Pro Ala Pro 385 390 395 400

Ala Lys Leu Cys Pro His 405

<210> 118

<211> 525

<212> PRT

<213> Homo sapien

<400> 118

Met Glu Ala Ala Ala Val Thr Val Thr Arg Ser Ala Thr Arg Arg Arg 1 5 10 15

Arg Arg Gln Leu Gln Gly Leu Ala Ala Pro Glu Ala Gly Thr Gln Glu  $20 \cdot 25$  30

Glu Gln Glu Asp Gln Glu Pro Arg Pro Arg Arg Arg Pro Gly Arg
35 40 45

Ser Ile Lys Asp Glu Glu Glu Glu Thr Val Phe Arg Glu Val Val Ser 50 55 60

Phe Ser Pro Asp Pro Leu Pro Val Arg Tyr Tyr Asp Lys Asp Thr Thr 65 70 75 80

Lys Pro Ile Ser Phe Tyr Leu Ser Ser Leu Glu Glu Leu Leu Ala Trp 85 90 95

Lys Pro Arg Leu Glu Asp Gly Phe Asn Val Ala Leu Glu Pro Leu Ala 100 105 110

Cys Arg Gln Pro Pro Leu Ser Ser Gln Arg Pro Arg Thr Leu Leu Cys 115 120 125 His Asp Met Met Gly Gly Tyr Leu Asp Asp Arg Phe Ile Gln Gly Ser 130 135 Val Val Gln Thr Pro Tyr Ala Phe Tyr His Trp Gln Cys Ile Asp Val 150 155 160 Phe Val Tyr Phe Ser His His Thr Val Thr Ile Pro Pro Val Gly Trp 170 175 Thr Asn Thr Ala His Arg His Gly Val Cys Val Leu Gly Thr Phe Ile 180 185 190 Thr Glu Trp Asn Glu Gly Gly Arg Leu Cys Glu Ala Phe Leu Ala Gly 195 200 205 Asp Glu Arg Ser Tyr Gln Ala Val Ala Asp Arg Leu Val Gln Ile Thr 210 215 220 Gln Phe Phe Arg Phe Asp Gly Trp Leu Ile Asn Ile Glu Asn Ser Leu 225 230 235 Ser Leu Ala Ala Val Gly Asn Met Pro Pro Phe Leu Arg Tyr Leu Thr 245 250 Thr Gln Leu His Arg Gln Val Pro Gly Gly Leu Val Leu Trp Tyr Asp 260 265 270 Ser Val Val Gln Ser Gly Gln Leu Lys Trp Gln Asp Glu Leu Asn Gln 275 280 285 His Asn Arg Val Phe Phe Asp Ser Cys Asp Gly Phe Phe Thr Asn Tyr 290 295 300 Asn Trp Arg Glu Glu His Leu Glu Arg Met Leu Gly Gln Ala Gly Glu 305 310 315 Arg Arg Ala Asp Val Tyr Val Gly Val Asp Val Phe Ala Arg Gly Asn 325 330 335 Val Val Gly Gly Arg Phe Asp Thr Asp Lys Ser Leu Glu Leu Ile Arg 340 345 350

Lys His Gly Phe Ser Val Ala Leu Phe Ala Pro Gly Trp Val Tyr Glu 355 360 365

Cys Leu Glu Lys Lys Asp Phe Phe Gln Asn Gln Asp Lys Phe Trp Gly 370 375 380

Arg Leu Glu Arg Tyr Leu Pro Thr His Ser Ile Cys Ser Leu Pro Phe 385 390 395 400

Val Thr Ser Phe Cys Leu Gly Met Gly Ala Arg Arg Val Cys Tyr Gly 405 410 415

Gln Glu Glu Ala Val Gly Pro Trp Tyr His Leu Ser Ala Gln Glu Ile 420 425 430

Gln Pro Leu Phe Gly Glu His Arg Leu Gly Gly Asp Gly Arg Gly Trp  $435 \ \ 440 \ \ 445$ 

Val Arg Thr His Cys Cys Leu Glu Asp Ala Trp His Gly Gly Ser Ser 450 455 460

Leu Leu Val Arg Gly Val Ile Pro Pro Glu Val Gly Asn Val Ala Val
465 470 475 480

Arg Trp Val Ser Asp Gly Gly Arg Trp Ala His Gln Leu Leu Pro Ser 485 490 495

His Val Val Ala Met Glu Leu Asp Arg Trp Gly Ser Gly Gln Asn 500 505 510

Lys Asp Arg Gly Gln Thr Gln Met Gly Phe Leu Lys Leu 515 520

<210> 119

<211> 327

<212> PRT

<213> Homo sapien

<400> 119

Met Phe Gln Pro Thr Lys Glu Ser Gly Leu Gly Gly Gly Leu Val Pro 1 5 10 15

Trp Leu Arg Thr Gly Pro Arg Cys Gly Ser Ile Trp His Pro Gly Pro 20 25 30

Leu Phe Leu Glu Gly Gly Ala Gly Gly Arg Asp Leu Glu Leu Ala Ser

171

35 40 45

Ile Ser Gln Trp Ser Leu His Gly Thr His His Arg Thr Phe Phe Pro 50 55 60

Arg Leu Phe Ser Leu Gln Ala Pro Val Pro Pro Lys Ile Tyr Leu Ser 65 70 75 80

Met Val Tyr Lys Leu Glu Gly Pro Thr Asp Val Thr Val Ala Leu Glu 85 90 95

Leu Thr Thr Gly Asp Ala Gly Ser Cys His Ile Gly Gly Ile Ser Val

Leu Asn Ala Glu Thr Ser Ser Arg His Ser Leu Arg Pro Leu Arg Val

Pro Pro Thr Lys Leu Ala Arg Trp Val Gly Arg Cys Gly Arg Gln Leu 130 135 140

Ser Gly Gly Trp Val Gln His Cys Tyr Glu Val Ser Leu Arg Gly Cys 145 150 155 160

Leu Leu Asp Leu Leu Val Cys Phe Ser Arg Pro Pro Gly Ser Arg

Glu Glu Glu Ser Phe Thr Cys Arg Leu Gly Glu Ile Gln Val Val Asp 180 185 190

Ala Ala Ser Leu Leu Ala Pro Leu Pro Gln Val Gln Ala Val Thr Ile 195 200 205

Ser His Ile Arg Trp Gln Pro Ser Ala Ser Glu Arg Glu Gly Pro Pro 210 215 220

Ala Leu Leu Gln Leu Ser Cys Thr Leu His Trp Ser Phe Leu Leu Ser 225 230 235 240

Gln Val Arg Cys Phe Arg Ile His Cys Trp Gly Gly Met Ser Asp Asp 245 250 255

Ser Pro Gly Arg Glu Leu Pro Arg Pro Glu Met Pro Met Phe Leu Gly 260 265 270

Leu Ala Phe Ala Thr Gln Tyr Arg Ile Val Asp Leu Leu Val Glu Ala 275 280 285

Ala Gly Pro Gly Gln Asp Arg Arg Met Glu Phe Leu Val Glu Pro Val 290 295 300

Pro Lys Glu Gly Phe Arg Val Pro Gln Ala Glu Trp Gly Arg Ala Val 305 310 315 320

Leu Leu Tyr Ser Ala Pro Ala 325

<210> 120

<211> 384

<212> PRT

<213> Homo sapien

<400> 120

Gln Ile Pro Arg Thr Val Ser Ser Cys Arg Thr Gly Leu Ser Pro Leu 1 5 10 15

His Ile Ser Pro Pro Ser Ser Pro Ser Pro Pro Lys Pro Pro Leu Tyr 20 25 30

Ser Ala Ser Val Ser Leu Asp Thr Leu Asp Ala Pro Tyr Glu Gly Ile 35 40 45

Pro Tyr Gly Ile Ser Glu Leu Arg Cys Phe Ser Pro Gln Lys Asn Leu 50 55 60

Ala Leu Gly Glu Asp Leu Ser Pro Gly Tyr Gly Gln Asp His Asp Val 65 70 75 80

Gly Ala Phe Gly Thr Gln Ala Pro Cys Ser Trp Arg Glu Gly Leu Val 85 90 95

Asp Ala Ile Trp Ser Trp Leu Arg Phe Leu Ser Gly Leu Ser Thr Ala

Pro Ile Thr Gly Pro Phe Ser Pro Gly Tyr Phe Tyr Ser Leu Gln Ala 115 120 125

Pro Val Pro Pro Lys Ile Tyr Leu Ser Met Val Tyr Lys Leu Glu Gly 130 135 140

Pro Thr Asp Val Thr Val Ala Leu Glu Leu Thr Thr Gly Asp Ala Gly 145 150 155 160

173

Ser Cys His Ile Gly Gly Ile Ser Val Leu Asn Ala Glu Thr Ser Ser 165 170 175

Arg His Ser Leu Arg Pro Leu Arg Val Pro Pro Thr Lys Leu Ala Arg 180 185 190

Trp Val Gly Arg Cys Gly Arg Gln Leu Ser Gly Gly Trp Val Gln His
195 200 205

Cys Tyr Glu Val Ser Leu Arg Gly Cys Leu Leu Leu Asp Leu Leu Val 210 215 220

Cys Phe Ser Arg Pro Pro Gly Ser Arg Glu Glu Glu Ser Phe Thr Cys 225 230 235 240

Arg Leu Gly Glu Ile Gin Val Val Asp Ala Ala Ser Leu Leu Ala Pro 245 250 255

Leu Pro Gln Val Gln Ala Val Thr Ile Ser His Ile Arg Trp Gln Pro 260 265 270

Ser Ala Ser Glu Arg Glu Gly Pro Pro Ala Leu Leu Gln Leu Ser Cys 275 280 285

Thr Leu His Trp Ser Phe Leu Leu Ser Gln Val Arg Cys Phe Arg Ile 290 295 300

His Cys Trp Gly Gly Met Ser Asp Asp Ser Pro Gly Arg Glu Leu Pro 305 310 315

Arg Pro Glu Met Pro Met Phe Leu Gly Leu Ala Phe Ala Thr Gln Tyr 325 330 335

Arg Ile Val Asp Leu Leu Val Glu Ala Ala Gly Pro Gly Gln Asp Arg 340 345 350

Arg Met Glu Phe Leu Val Glu Pro Val Pro Lys Glu Gly Phe Arg Val 355 360 365

Pro Gln Ala Glu Trp Gly Arg Ala Val Leu Leu Tyr Ser Ala Pro Ala 370 380

<210> 121

<211> 450

<212> PRT

The state of the s

<213> Homo sapien

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| 1 |   |
|---|---|
|   | 1 |

Gln Ile Pro Arg Thr Val Ser Ser Cys Arg Thr Gly Leu Ser Pro Leu 1 5 10 15

His Ile Ser Pro Pro Ser Ser Pro Ser Pro Pro Lys Pro Pro Leu Tyr
20 25 30

Ser Ala Ser Val Ser Leu Asp Thr Leu Asp Ala Pro Tyr Glu Gly Ile 35 40 45

Pro Tyr Gly Ile Ser Glu Leu Arg Cys Phe Ser Pro Gln Lys Asn Leu 50 55 60

Ala Leu Gly Glu Asp Leu Ser Pro Gly Tyr Gly Gln Asp His Asp Val 65 70 75 80

Gly Ala Phe Gly Thr Gln Ala Pro Cys Ser Trp Arg Glu Gly Leu Val 85 90 95

Asp Ala Ile Trp Ser Trp Leu Arg Phe Leu Ser Gly Leu Ser Thr Ala 100 105 110

Pro Ile Thr Gly Pro Phe Ser Pro Gly Tyr Phe Tyr Ser Leu Gln Ala 115 120 125

Pro Val Pro Pro Lys Ile Tyr Leu Ser Met Val Tyr Lys Leu Glu Gly 130 135 140

Pro Thr Asp Val Thr Val Ala Leu Glu Leu Thr Thr Gly Asp Ala Gly 145 150 155

Ser Cys His Ile Gly Gly Ile Ser Val Leu Asn Ala Glu Thr Ser Ser 165 170 175

Arg His Ser Leu Arg Pro Leu Arg Val Pro Pro Thr Lys Leu Ala Arg 180 185 190

Trp Val Gly Arg Cys Gly Arg Gln Leu Ser Gly Gly Trp Val Gln His
195 200 205

Cys Tyr Glu Val Ser Leu Arg Gly Cys Leu Leu Leu Asp Leu Leu Val 210 215 220

Cys Phe Ser Arg Pro Pro Gly Ser Arg Glu Glu Glu Ser Phe Thr Cys 225 230 235 240

175

Arg Leu Gly Glu Ile Gln Val Met Leu Pro Arg Gly Ala Arg Ala Gly 245 250 255

Leu Ala Val Cys Pro Ala Gly Val Gly Val Glu Ala Ala Pro Gly Arg
260 265 270

Pro Leu Leu Gly Phe Ser Gly Glu Leu Gly Trp Arg Ser Gln Gly Gly 275 280 285

Glu Met Cys Ala Trp Gly His Pro Leu Pro Ala Pro Gly Arg Pro Ala 290 295 300

Val Leu Ser Leu Leu Ser Cys Gln Val Val Asp Ala Ala Ser Leu Leu 305 310 315 320

Ala Pro Leu Pro Gln Val Gln Ala Val Thr Ile Ser His Ile Arg Trp
325 330 335

Gln Pro Ser Ala Ser Glu Arg Glu Gly Pro Pro Ala Leu Leu Gln Leu 340 345 350

Ser Cys Thr Leu His Trp Ser Phe Leu Leu Ser Gln Val Arg Cys Phe 355 360 365

Arg Ile His Cys Trp Gly Gly Met Ser Asp Asp Ser Pro Gly Arg Glu 370 380

Leu Pro Arg Pro Glu Met Pro Met Phe Leu Gly Leu Ala Phe Ala Thr 385 390 395 400

Gln Tyr Arg Ile Val Asp Leu Leu Val Glu Ala Ala Gly Pro Gly Gln 405 410 415

Asp Arg Arg Met Glu Phe Leu Val Glu Pro Val Pro Lys Glu Gly Phe 420 425 430

Arg Val Pro Gln Ala Glu Trp Gly Arg Ala Val Leu Leu Tyr Ser Ala 435 440 445

Pro Ala 450

<210> 122

<211> 302

<212> PRT

<213> Homo sapien

| e i | A 1 | ٠, | ٦ | _ | 1 | 2 | 2 |
|-----|-----|----|---|---|---|---|---|
| ~ ' | 41  |    |   | , | _ | ~ | 4 |

Met Glu Ala Ala Val Thr Val Thr Arg Ser Ala Thr Arg Arg Arg 1 5 10 15

Arg Arg Gln Leu Gln Gly Leu Ala Ala Pro Glu Ala Gly Thr Gln Glu 20 25 30

Glu Gln Glu Asp Gln Glu Pro Arg Pro Arg Arg Arg Pro Gly Arg 35 40 45

Ser Ile Lys Asp Glu Glu Glu Glu Thr Val Phe Arg Glu Val Val Ser 50 55

Phe Ser Pro Asp Pro Leu Pro Val Arg Tyr Tyr Asp Lys Asp Thr Thr 65 70 75 80

Lys Pro Ile Ser Phe Tyr Leu Ser Ser Leu Glu Glu Leu Leu Ala Trp 85 90 95

Lys Pro Arg Leu Glu Asp Gly Phe Asn Val Ala Leu Glu Pro Leu Ala 100 105 110

Cys Arg Gln Pro Pro Leu Ser Ser Gln Arg Pro Arg Thr Leu Leu Cys 115 120 125

His Asp Met Met Gly Gly Tyr Leu Asp Asp Arg Phe Ile Gln Gly Ser 130 135 140

Val Val Gln Thr Pro Tyr Ala Phe Tyr His Trp Gln Cys Ile Asp Val 145 150 155 160

Phe Val Tyr Phe Ser His His Thr Val Thr Ile Pro Pro Val Gly Trp 165 170 175

Thr Asn Thr Ala His Arg His Gly Val Cys Val Leu Gly Thr Phe Ile 180 185 190

Thr Glu Trp Asn Glu Gly Gly Arg Leu Cys Glu Ala Phe Leu Ala Gly 195 200 205

Asp Glu Arg Ser Tyr Gln Ala Val Ala Asp Arg Leu Val Gln Ile Thr 210 215 220

Gln Phe Phe Arg Phe Asp Gly Trp Leu Ile Asn Ile Glu Asn Ser Leu 225 230 235 240

Ser Leu Ala Ala Val Gly Asn Met Pro Pro Phe Leu Arg Tyr Leu Thr 245 250 255

Thr Gln Leu Leu Val Glu Ala Ala Gly Pro Gly Gln Asp Arg Met 260 265 270

Glu Phe Leu Val Glu Pro Val Pro Lys Glu Gly Phe Arg Val Pro Gln 275 280 285

Ala Glu Trp Gly Arg Ala Val Leu Leu Tyr Ser Ala Pro Ala 290 295 300

<210> 123

<211> 162

<212> PRT

<213> Homo sapien

<400> 123

Met Pro Arg Trp Tyr Phe Leu Leu Ala Arg Cys Phe Gly Cys Ala Val 1 5 10 15

Ile Glu Asp Thr Trp His Tyr Phe Leu His Arg Leu Leu His His Lys 20 25 30

Arg Ile Tyr Lys Tyr Ile His Lys Val His His Glu Phe Gln Ala Pro 35 40 45

Phe Gly Met Glu Ala Glu Tyr Ala His Pro Leu Glu Thr Leu Ile Leu 50 55 60

Gly Thr Gly Phe Phe Ile Gly Ile Val Leu Leu Cys Asp His Val Ile 65 70 75 80

Leu Leu Trp Ala Trp Val Thr Ile Arg Leu Leu Glu Thr Ile Asp Val 85 90 95

His Ser Gly Tyr Asp Ile Pro Leu Asn Pro Leu Asn Leu Ile Pro Phe 100 105 110

Tyr Ala Gly Ser Arg His His Asp Phe His His Met Asn Phe Ile Gly 115 120 125

Asn Tyr Ala Ser Thr Phe Thr Trp Trp Asp Arg Ile Phe Gly Thr Asp 130 135 140

178

Ser Gln Tyr Asn Ala Tyr Asn Glu Lys Arg Lys Lys Phe Glu Lys Lys 145 150 155 160

Thr Glu

<210> 124

<211> 206

<212> PRT

<213> Homo sapien

<400> 124

Met Gly Glu Pro Gln Gly Ser Met Arg Ile Leu Val Thr Gly Gly Ser 1 5 10 15

Gly Leu Val Gly Lys Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly 20 25 30

Leu Pro Gly Glu Asp Trp Val Phe Val Ser Ser Lys Asp Ala Asp Leu 35 40 45

Thr Asp Thr Ala Gln Thr Arg Ala Leu Phe Glu Lys Val Gln Pro Thr 50 . 55 60

His Val Ile His Leu Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile 65 70 75 80

Lys Tyr Asn Leu Asp Phe Trp Arg Lys Asn Val His Met Asn Asp Asn 85 90 95

Val Leu His Ser Ala Phe Glu Val Gly Ala Arg Lys Val Val Ser Cys 100 105 110

Leu Ser Thr Cys Ile Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu 115 120 125

Thr Met Ile His Asn Gly Pro Pro His Asn Ser Asn Phe Gly Tyr Ser 130 135 140

Tyr Ala Lys Arg Met Ile Asp Val Gln Asn Ser Ala Gly Pro Thr Ser 145 150 155 160

Ser Ser Thr Ala Ala Pro Ser Pro Leu Ser Ser Pro Pro Thr Ser Ser 165 170 175

Gly Pro Thr Thr Ser Thr Ser Arg Met Ala Thr Cys Cys Leu Ala 180 185 190

Ser Ser Thr Arg Cys Thr Trp Pro Arg Ala Ala Ala Arg Pro 200

<210> 125 <211> 380

<212> PRT

<213> Homo sapien

<400> 125

Leu Gln Val Pro Ala Val Pro Gly Thr Leu Arg Ala Pro Gly Thr Pro

Phe Pro Arg Val Pro Arg Pro Ser Leu Pro Ala Pro Pro Pro Thr Trp

Leu Arg Gly Gln Pro Glu Arg Thr Arg Pro Glu Ala Ala Val Gly Glu 40

Pro Ala Val Gly Leu Asp Ala Gly Ala Thr Asp Met Gly Glu Pro Gln 55

Gly Ser Met Arg Ile Leu Val Thr Gly Gly Ser Gly Leu Val Gly Lys

Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly Leu Pro Gly Glu Asp 85

Trp Val Phe Val Ser Ser Lys Asp Ala Asp Leu Thr Asp Thr Ala Gln 110

Thr Arg Ala Leu Phe Glu Lys Val Gln Pro Thr His Val Ile His Leu 120

Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile Lys Tyr Asn Leu Asp

Phe Trp Arg Lys Asn Val His Met Asn Asp Asn Val Leu His Ser Ala 150 155

Phe Glu Val Gly Ala Arg Lys Val Val Ser Cys Leu Ser Thr Cys Ile 165 170

Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu Thr Met Ile His Asn 185

Gly Pro Pro His Asn Ser Asn Phe Gly Tyr Ser Tyr Ala Lys Arg Met 195 200 205 .

Ile Asp Val Gln Asn Arg Ala Tyr Phe Gln Gln Tyr Gly Cys Thr Phe 210 215 220

Thr Ala Val Ile Pro Thr Asn Val Phe Gly Pro His Asp Asn Phe Asn 225 230 235 240

Ile Glu Asp Gly His Val Leu Pro Gly Leu Ile His Lys Val His Leu 245 250 255

Ala Lys Ser Ser Gly Ser Ala Leu Thr Val Trp Gly Thr Gly Asn Pro 260 265 270

Arg Arg Gln Phe Ile Tyr Ser Leu Asp Leu Ala Gln Leu Phe Ile Trp 275 280 285

Val Leu Arg Glu Tyr Asn Glu Val Glu Pro Ile Ile Leu Ser Val Gly
290 295 300

Glu Glu Asp Glu Val Ser Ile Lys Glu Ala Ala Glu Ala Val Val Glu 305 310 315 320

Ala Met Asp Phe His Gly Glu Val Thr Phe Asp Thr Thr Lys Ser Asp 325 330 335

Gly Gln Phe Lys Lys Thr Ala Ser Asn Ser Lys Leu Arg Thr Tyr Leu 340 345 350

Pro Asp Phe Arg Phe Thr Pro Phe Lys Gln Ala Val Lys Glu Thr Cys 355 360 365

Ala Trp Phe Thr Asp Asn Tyr Glu Gln Ala Arg Lys 370 375 380

<210> 126

<211> 380

<212> PRT

<213> Homo sapien

<400> 126

Leu Gln Val Pro Ala Val Pro Gly Thr Leu Arg Ala Pro Gly Thr Pro 1 5 10 15

Phe Pro Arg Val Pro Arg Pro Ser Leu Pro Ala Pro Pro Pro Thr Trp 20 25 30

Leu Arg Gly Gln Pro Glu Arg Thr Arg Pro Glu Ala Ala Val Gly Glu 35 40 45

Pro Ala Val Gly Leu Asp Ala Gly Ala Thr Asp Met Gly Glu Pro Gln 50 55 60

Gly Ser Met Arg Ile Leu Val Thr Gly Gly Ser Gly Leu Val Gly Lys 65 70 75 80

Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly Leu Pro Gly Glu Asp 85 90 95

Thr Arg Ala Leu Phe Glu Lys Val Gln Pro Thr His Val Ile His Leu 115 120 125

Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile Lys Tyr Asn Leu Asp 130 135 140

Phe Trp Arg Lys Asn Val His Met Asn Asp Asn Val Leu His Ser Ala 145 150 155 160

Phe Glu Val Gly Ala Arg Lys Val Val Ser Cys Leu Ser Thr Cys Ile 165 170 175

Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu Thr Met Ile His Asr 180 185 190

Gly Pro Pro His Asn Ser Asn Phe Gly Tyr Ser Tyr Ala Lys Arg Met 195 200 205

Ile Asp Val Gln Asn Arg Ala Tyr Phe Gln Gln Tyr Gly Cys Thr Phe 210 215 220

Thr Ala Val Ile Pro Thr Asn Val Phe Gly Pro His Asp Asn Phe Asn 225 230 235 240

Ile Glu Asp Gly His Val Leu Pro Gly Leu Ile His Lys Val His Leu 245 250 255

Ala Lys Ser Ser Gly Ser Ala Leu Thr Val Trp Gly Thr Gly Asn Pro - 260 265 270

Arg Arg Gln Phe Ile Tyr Ser Leu Asp Leu Ala Gln Leu Phe Ile Trp 275

Val Leu Arg Glu Tyr Asn Glu Val Glu Pro Ile Ile Leu Ser Val Gly 295 290

Glu Glu Asp Glu Val Ser Ile Lys Glu Ala Ala Glu Ala Val Val Glu 310

Ala Met Asp Phe His Gly Glu Val Thr Phe Asp Thr Thr Lys Ser Asp

Gly Gln Phe Lys Lys Thr Ala Ser Asn Ser Lys Leu Arg Thr Tyr Leu

Pro Asp Phe Arg Phe Thr Pro Phe Lys Gln Ala Val Lys Glu Thr Cys 360

Ala Trp Phe Thr Asp Asn Tyr Glu Gln Ala Arg Lys 375

<210> 127 <211> 334 <212> PRT <213> Homo sapien

<400> 127

Met Arg Ala Leu Ala Ala Asn Arg Val Asn Asp Leu Cys Gln Glu Pro 10

Pro Ser Gln Gly Cys Leu Pro Pro Pro Leu Val Ser Gln Arg Gly Val 25

Glu Cys Thr Phe Ser Arg Pro Ser Gly Glu Ser Trp Val Gly Thr Ser 40

Cys Ser Gly Leu Gly Gly Ser Ser Gly Pro Leu Arg Arg Cys Arg Leu 60

Arg Ala Pro Arg Gly Thr Gly Leu Arg Arg Gly Ser Ala Ser Val Gln 75 70

Leu Gly Leu Ser Gly Cys Gln Trp Thr Met Pro His Ser Glu Gly Leu 90 95 85

Thr Leu Cys Gln Leu Pro Gln Lys Ser Gly Ala Pro Lys Asp Glu Ser

183

100 105 110

Val Met Thr Ser Ala Ser His Cys Leu Thr Leu Gln Gly Ala Thr Asp 115 120 125

Met Gly Glu Pro Gln Gly Ser Met Arg Ile Leu Val Thr Gly Gly Ser 130 135 140

Gly Leu Val Gly Lys Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly 145 150 155 160

Leu Pro Gly Glu Asp Trp Val Phe Val Ser Ser Lys Asp Ala Asp Leu 165 170 175

Thr Asp Thr Ala Gln Thr Arg Ala Leu Phe Glu Lys Val Gln Pro Thr 180 185 190

His Val Ile His Leu Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile 195 200 205

Lys Tyr Asn Leu Asp Phe Trp Arg Lys Asn Val His Met Asn Asp Asn 210 215 220

Val Leu His Ser Ala Phe Glu Val Gly Ala Arg Lys Val Val Ser Cys 235 230 240

Leu Ser Thr Cys Ile Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu 245 250 255

Thr Met Ile His Asn Gly Pro Pro His Asn Ser Asn Phe Gly Tyr Ser 260 265 270

Tyr Ala Lys Arg Met Ile Asp Val Gln Asn Ser Ala Gly Pro Thr Ser 275 280 285

Ser Ser Thr Ala Ala Pro Ser Pro Leu Ser Ser Pro Pro Thr Ser Ser 290 295 300

Gly Pro Thr Thr Thr Ser Thr Ser Arg Met Ala Thr Cys Cys Leu Ala 305 310 315 320

Ser Ser Thr Arg Cys Thr Trp Pro Arg Ala Ala Arg Pro 325 330

<210> 128 <211> 327

<212> PRT

<213> Homo sapien

<400> 128

His Tyr Ser Ala Thr Asp Met Gly Glu Pro Gln Gly Ser Met Arg Ile 1 5 10 15

Leu Val Thr Gly Gly Ser Gly Leu Val Gly Lys Ala Ile Gln Lys Val 20 25 30

Val Ala Asp Gly Ala Gly Leu Pro Gly Glu Asp Trp Val Phe Val Ser 35 40 45

Ser Lys Asp Ala Asp Leu Thr Asp Thr Ala Gln Thr Arg Ala Leu Phe 50 60

Glu Lys Val Gln Pro Thr His Val Ile His Leu Ala Ala Met Val Gly 70 75 80

Gly Leu Phe Arg Asn Ile Lys Tyr Asn Leu Asp Phe Trp Arg Lys Asn 85 90 95

Val His Met Asn Asp Asn Val Leu His Ser Ala Phe Glu Val Gly Ala 100 105 110

Arg Lys Val Val Ser Cys Leu Ser Thr Cys Ile Phe Pro Asp Lys Thr 115 120 125

Thr Tyr Pro Ile Asp Glu Thr Met Ile His Asn Gly Pro Pro His Asn 130 135 140

Ser Asn Phe Gly Tyr Ser Tyr Ala Lys Arg Met Ile Asp Val Gln Asn 145 150 155 160

Arg Ala Tyr Phe Gln Gln Tyr Gly Cys Thr Phe Thr Ala Val Ile Pro 165 170 175

Thr Asn Val Phe Gly Pro His Asp Asn Phe Asn Ile Glu Asp Gly His 180 185 190

Val Leu Pro Gly Leu Ile His Lys Val His Leu Ala Lys Ser Ser Gly 195 200 205

Ser Ala Leu Thr Val Trp Gly Thr Gly Asn Pro Arg Arg Gln Phe Ile 210 215 220

185

Tyr Ser Leu Asp Leu Ala Gln Leu Phe Ile Trp Val Leu Arg Glu Tyr 225 230 235 240

Asn Glu Val Glu Pro Ile Ile Leu Ser Val Gly Glu Glu Asp Glu Val 245 250 255

Ser Ile Lys Glu Ala Ala Glu Ala Val Val Glu Ala Met Asp Phe His 260 265 270

Gly Glu Val Thr Phe Asp Thr Thr Lys Ser Asp Gly Gln Phe Lys Lys 275 280 285

Thr Ala Ser Asn Ser Lys Leu Arg Thr Tyr Leu Pro Asp Phe Arg Phe 290 295 300

Thr Pro Phe Lys Gln Ala Val Lys Glu Thr Cys Ala Trp Phe Thr Asp 305 310 315 320

Asn Tyr Glu Gln Ala Arg Lys 325

<210> 129

<211> 151

<212> PRT

<213> Homo sapien

<400> 129

Met Gly Glu Pro Gln Gly Ser Met Arg Ile Leu Val Thr Gly Gly Ser 1 5 10 15

Gly Leu Val Gly Lys Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly 20 25 30

Leu Pro Gly Glu Asp Trp Val Phe Val Ser Ser Lys Asp Ala Asp Leu 35 40 45

Thr Asp Thr Ala Gln Thr Arg Ala Leu Phe Glu Lys Val Gln Pro Thr 50 55 60

His Val Ile His Leu Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile 65 70 75 80

Lys Tyr Asn Leu Asp Phe Trp Arg Lys Asn Val His Met Asn Asp Asn 85 90 95

Val Leu His Ser Ala Phe Glu Val Gly Ala Arg Lys Val Val Ser Cys 100 105 110

Leu Ser Thr Cys Ile Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu 115 120 125

Thr Met Ile His Asn Gly Pro Pro His Asn Ser Asn Phe Gly Tyr Ser 130 135 140

Tyr Ala Lys Arg Met Ile Asp Val Gln Asn Arg Ser Ser Arg Pro Ser 145 150 155

Cys

<210> 130

<211> 326

<212> PRT

<213> Homo sapien

<400> 130

Leu Gln Val Pro Ala Val Pro Gly Thr Leu Arg Ala Pro Gly Thr Pro 1 5 10 15

Phe Pro Arg Val Pro Arg Pro Ser Leu Pro Ala Pro Pro Pro Thr Trp 20 25 30

Leu Arg Gly Gln Pro Glu Arg Thr Arg Pro Glu Ala Ala Val Gly Glu 35 40 45

Pro Ala Val Gly Leu Asp Ala Gly Ala Thr Asp Met Gly Glu Pro Gln 50 55 60

Gly Ser Met Arg Ile Leu Val Thr Gly Gly Ser Gly Leu Val Gly Lys 70 75 80

Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly Leu Pro Gly Glu Asp 85 90 95

Trp Val Phe Val Ser Ser Lys Asp Ala Asp Leu Thr Asp Thr Ala Gln
100 105 110

Thr Arg Ala Leu Phe Glu Lys Val Gln Pro Thr His Val Ile His Leu 115 120 125

Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile Lys Tyr Asn Leu Asp 130 135 140

187

Phe Trp Arg Lys Asn Val His Met Asn Asp Asn Val Leu His Ser Ala 145 150 155 160

Phe Glu Val Gly Ala Arg Lys Val Val Ser Cys Leu Ser Thr Cys Ile 165 170 175

Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu Thr Met Ile His Asn 180 185 190

Gly Pro Pro His Asn Ser Asn Phe Gly Tyr Ser Tyr Ala Lys Arg Met 195 200 205

Ile Asp Val Gln Asn Arg Ala Tyr Phe Gln Gln Tyr Gly Cys Thr Phe 210 225 220

Thr Ala Val Ile Pro Thr Asn Val Phe Gly Pro His Asp Asn Phe Asn 225 230 235 240

Ile Glu Asp Gly His Val Leu Pro Gly Leu Ile His Lys Val His Leu 245  $\phantom{0}$  250  $\phantom{0}$  255

Ala Lys Ser Ser Gly Ser Ala Leu Thr Val Trp Gly Thr Gly Asn Pro 260 265 270

Arg Arg Gln Phe Ile Tyr Ser Leu Asp Leu Ala Gln Leu Phe Ile Trp 275 280 285

Val Leu Arg Glu Tyr Asn Glu Val Glu Pro Ile Ile Leu Ser Gly Gly 290 295 300

Tyr Leu Ser Pro Gln Pro Pro Ser Ser Met Val Gly Gln Asp Pro Arg 305 310 315 320

Leu Ser Trp Glu Ala Gly 325

<210> 131

<211> 216

<212> PRT

<213> Homo sapien

<400> 131

Met Gln Val Arg Thr Asp Pro Arg Ser Arg Gln Cys Trp Pro Leu Glu 1 5 10 15

His Arg Thr Trp Leu Thr Asp Ser His Ser Ser Cys Leu Phe Pro Leu 20 25 30

Pro Leu Glu Gln Pro Ser Leu Leu Gln Ser Asn Pro Cys Pro Ser Phe 35 40 45

Leu Pro Leu Ser Arg Ala Ala Pro Pro Ala His Leu Arg Pro Gly Pro 55

Ser Tyr Leu Leu Pro Leu Leu Ser Cys Pro Ile Pro Val Val Arg Arg 65 70 75

Glu Ser Thr Gly Gln Arg Pro Ser Ser Thr Cys Asp Leu Gly Glu Cys 
 85
 90
 95

Gln Ala Ser Pro Arg Gly Pro Gly Pro Arg Gly Pro Gly Arg Leu Cys 100 105 110

Cys Gly Gly Ser Arg Val Arg Thr Gly Ala Ala Ser Pro Leu Ala Val 115 120 125

Cys Leu Cys Pro Leu His Trp Pro Leu Glu Ala Gln Arg Pro Ser Glv 130 135

Tyr Phe Pro Ser Ser Gly Leu Pro Leu Met Leu Phe Pro Ala Pro Phe 145 150 155 160

Phe Tyr Leu Glu Thr Pro Ile Pro Ser His Pro Leu Gln Arg Ser Ser 165 170 175

Gln Ser Cys Pro Gln His Gly Ser Leu His Ser Pro Trp Val Ser Pro 180 185 190

Pro Val Ser Cys Leu Pro Arg Thr Pro Asp Leu Pro Leu Pro Gly Trp 195 200 205

Pro Arg Trp Ile Leu Tyr Ser Asp 210 215

<210> 132 <211> 108 <212> PRT <213> Homo sapien

<400> 132

Met Ala His Ala Thr Leu Ser Ala Ala Pro Ser Asn Pro Arg Leu Leu

189

Arg Val Ala Leu Leu Leu Leu Leu Val Ala Ala Ser Arg Arg Ala 20 25 30

Ala Gly Gly Ser Arg Arg Pro Gly Val Pro Gly Pro Asp Ala Ala Gly 35  $\phantom{\bigg|}40\phantom{\bigg|}40\phantom{\bigg|}45\phantom{\bigg|}$ 

Val Gly Ala Pro Arg Arg Thr Ala Pro Leu Asn Gln Arg Val Tyr Ser 50 55

Ser Leu Gly Ala Ser Val Val Thr Glu Leu Arg Cys Gln Cys Leu Gln 65 70 75 80

Thr Leu Gln Gly Ile His Leu Lys Asn Ile Gln Ser Val Asn Val Arg 85 90 95

Ser Pro Gly Pro His Cys Ala Gln Thr Glu Val Met 100 105

<210> 133

<211> 142

<212> PRT

<213> Homo sapien

<400> 133

Lys Gly Ser Pro Ile Leu Gly Ser His Thr Ala Arg Val Ala Gly Thr 1 5 10 15

Ser Pro Pro Ala Leu Pro Leu Leu Ala Gln Leu Pro Asp Ala Ser Ala 20 25 30

Glu Pro His Gly Pro Arg His Ala Leu Arg Arg Pro Gln Gln Ser Pro 35 40 45

Ala Pro Ala Gly Gly Ala Ala Ala Pro Ala Pro Gly Gly Arg Gln Pro 50 60

Ala Arg Ser Arg Trp Val Pro Ala Pro Trp Gly Pro Arg Ala Gly Arg 65 70 75 80

Gly Trp Gly Gly Arg Pro Ala Pro Thr Ala Pro Leu Asn Gln Arg Val 85 90 95

Tyr Ser Ser Leu Gly Ala Ser Val Val Thr Glu Leu Arg Cys Gln Cys 100 105 110

Leu Gln Thr Leu Gln Gly Ile His Leu Lys Asn Ile Gln Ser Val Asn

Val Arg Ser Pro Gly Pro His Cys Ala Gln Thr Glu Val Met 130 135

<210> 134

<211> 482 <212> PRT <213> Homo sapien

<400> 134

Met Val Met Glu Lys Pro Ser Pro Leu Leu Val Gly Arg Glu Phe Val

Arg Gln Tyr Tyr Thr Leu Leu Asn Lys Ala Pro Glu Tyr Leu His Arg 25

Phe Tyr Gly Arg Asn Ser Ser Tyr Val His Gly Gly Val Asp Ala Ser 40

Gly Lys Pro Gln Glu Ala Val Tyr Gly Gln Asn Asp Ile His His Lys 50

Val Leu Ser Leu Asn Phe Ser Glu Cys His Thr Lys Ile Arg His Val

Asp Ala His Ala Thr Leu Ser Asp Gly Val Val Val Gln Val Met Gly 90

Leu Leu Ser Asn Ser Gly Gln Pro Glu Arg Lys Phe Met Gln Thr Phe 105

Val Leu Ala Pro Glu Gly Ser Val Pro Asn Lys Phe Tyr Val His Asn

Asp Met Phe Arg Tyr Glu Asp Glu Val Phe Gly Asp Ser Glu Pro Glu

Leu Asp Glu Glu Ser Glu Asp Glu Val Glu Glu Glu Glu Glu Arg 150 155

Gln Pro Ser Pro Glu Pro Val Gln Glu Asn Ala Asn Ser Gly Tyr Tyr 165 170

Glu Ala His Pro Val Thr Asn Gly Ile Glu Glu Pro Leu Glu Glu Ser 185 180

191

Ser His Glu Pro Glu Pro Glu Pro Glu Ser Glu Thr Lys Thr Glu Glu
195 200 205

Leu Lys Pro Gln Val Glu Glu Lys Asn Leu Glu Glu Leu Glu Glu Lys 210 215 220

Ser Thr Thr Pro Pro Pro Ala Glu Pro Val Ser Leu Pro Gln Glu Pro 225 230 235 240

Pro Lys Ala Phe Ser Trp Ala Ser Val Thr Ser Lys Asn Leu Pro Pro 245 250 255

Ser Gly Thr Val Ser Ser Ser Gly Ile His Pro Met Leu Lys His Gln 260 265 270

Ser His Ser Gln Glu Ser Lys Leu Asn Gln Lys Phe Asn Leu Ser His 275 280 285

Leu Val Cys Val Asn Asn Asp Leu Glu Asn Asp Leu Val Phe Leu Leu 290 295 300

Glu Asp Gln Asp Gln Ala Glu Glu Ile Met Glu Gln Asn Asp Ser Asp 305 310 315 320

Asn Arg Arg Ile Ile Arg Tyr Pro Asp Ser His Gln Leu Phe Val Gly 325 330 335

Asn Leu Pro His Asp Ile Asp Glu Asn Glu Leu Lys Glu Phe Phe Met 340 350

Ser Phe Gly Asn Val Val Glu Leu Arg Ile Asn Thr Lys Gly Val Gly 355 360 365

Gly Lys Leu Pro Asn Phe Gly Phe Val Val Phe Asp Asp Ser Glu Pro 370 375 380

Val Gln Arg Ile Leu Ile Ala Lys Pro Ile Met Phe Arg Gly Glu Val 385 390 395 400

Arg Leu Asn Val Glu Glu Lys Lys Thr Arg Ala Ala Arg Glu Arg Glu 415

Thr Arg Gly Gly Asp Asp Arg Arg Asp Ile Arg Arg Asn Asp Arg 420 425 430

Gly Pro Gly Gly Pro Arg Gly Ile Val Gly Gly Met Met Arg Asp

192

435 440 445

Arg Asp Gly Arg Gly Pro Pro Arg Gly Gly Met Ala Gln Lys Leu
450 455

Gly Ser Gly Arg Gly Thr Gly Gln Met Glu Gly Arg Phe Thr Gly Gln 465 470 475 480

Arg Arg

<210> 135

<211> 392

<212> PRT

<213> Homo sapien

<400> 135

Leu Ser Arg Ser Trp Val Cys Cys Leu Thr Val Asp Asn Gln Lys Glu 1 5 10 15

Ser Leu Cys Lys Pro Leu Phe Trp Leu Leu Lys Asp Leu Phe Gln Ile 20 25 30

Asn Phe Met Phe Thr Met Ile Cys Phe Val Met Lys Met Lys Cys Tyr 35 40 45

Gly Asp Ser Glu Pro Glu Leu Asp Glu Glu Ser Glu Asp Glu Val Glu 50 55 60

Glu Glu Glu Glu Glu Arg Gln Pro Ser Pro Glu Pro Val Gln Glu Asn 65 70 75 80

Ala Asn Ser Gly Tyr Tyr Glu Ala His Pro Val Thr Asn Gly Ile Glu 85 90 95

Glu Pro Leu Glu Glu Ser Ser His Glu Pro Glu Pro Glu Pro Glu Ser 100 105 110

Glu Thr Lys Thr Glu Glu Leu Lys Pro Gln Val Glu Glu Lys Asn Leu 115 120 125

Glu Glu Leu Glu Glu Lys Ser Thr Thr Pro Pro Pro Ala Glu Pro Val 130 140

Ser Leu Pro Gln Glu Pro Pro Lys Ala Phe Ser Trp Ala Ser Val Thr 145 150 155 160

193

Ser Lys Asn Leu Pro Pro Ser Gly Thr Val Ser Ser Ser Gly Ile Pro 165 170 175

Pro His Val Lys Ala Pro Val Ser Gln Pro Arg Val Glu Ala Lys Pro 180 185 190

Glu Val Gln Ser Gln Pro Pro Arg Val Arg Glu Gln Arg Pro Arg Glu 195 200 205

Arg Pro Gly Phe Pro Pro Arg Gly Pro Arg Pro Gly Arg Gly Asp Met 210 215 220

Glu Gln Asn Asp Ser Asp Asn Arg Arg Ile Ile Arg Tyr Pro Asp Ser 225 230 235 240

His Gln Leu Phe Val Gly Asn Leu Pro His Asp Ile Asp Glu Asn Glu 245 250 255

Leu Lys Glu Phe Phe Met Ser Phe Gly Asn Val Val Glu Leu Arg Ile 260 265 270

Asn Thr Lys Gly Val Gly Gly Lys Leu Pro Asn Phe Gly Phe Val Val 275 280 285

Phe Asp Asp Ser Glu Pro Val Gln Arg Ile Leu Ile Ala Lys Pro Ile 290 295 300

Met Phe Arg Gly Glu Val Arg Leu Asn Val Glu Glu Lys Lys Thr Arg 305 310 315 320

Ala Ala Arg Glu Arg Glu Thr Arg Gly Gly Gly Asp Asp Arg Arg Asp 325 330 335

Ile Arg Arg Asn Asp Arg Gly Pro Gly Gly Pro Arg Gly Ile Val Gly 340 345 350

Gly Gly Met Met Arg Asp Arg Gly Arg Gly Pro Pro Arg Gly 355 360 365

Gly Met Ala Gln Lys Leu Gly Ser Gly Arg Gly Thr Gly Gln Met Glu  $370 \hspace{1cm} 375 \hspace{1cm} 380$ 

Gly Arg Phe Thr Gly Gln Arg Arg 385 390

<210> 136

<211> 316

<212> PRT

<213> Homo sapien

<400> 136

Asp Trp Glu Glu Lys Arg Val Leu Ala Ile Cys Leu Ala Ser Gln Ser 1 5 10 15

Glu Asp Glu Val Glu Glu Glu Glu Glu Arg Gln Pro Ser Pro Glu 20 25 30

Pro Val Gln Glu Asn Ala Asn Ser Gly Tyr Tyr Glu Ala His Pro Val 35 40 45

Thr Asn Gly Ile Glu Glu Pro Leu Glu Glu Ser Ser His Glu Pro Glu 50 55 60

Pro Glu Pro Glu Ser Glu Thr Lys Thr Glu Glu Leu Lys Pro Gln Val 65 70 75 80

Glu Glu Lys Asn Leu Glu Glu Leu Glu Glu Lys Ser Thr Thr Pro Pro 85 90 95

Pro Ala Glu Pro Val Ser Leu Pro Gln Glu Pro Pro Lys Pro Arg Val

Glu Ala Lys Pro Glu Val Gln Ser Gln Pro Pro Arg Val Arg Glu Gln
115 120 125

Arg Pro Arg Glu Arg Pro Gly Phe Pro Pro Arg Gly Pro Arg Pro Gly 130 135 140

Arg Gly Asp Met Glu Gln Asn Asp Ser Asp Asn Arg Arg Ile Ile Arg 145 150 155 160

Tyr Pro Asp Ser His Gln Leu Phe Val Gly Asn Leu Pro His Asp Ile 165 170 175

Asp Glu Asn Glu Leu Lys Glu Phe Phe Met Ser Phe Gly Asn Val Val 180 180 185

Glu Leu Arg Ile Asn Thr Lys Gly Val Gly Gly Lys Leu Pro Asn Phe 195 200 205

Gly Phe Val Val Phe Asp Asp Ser Glu Pro Val Gln Arg Ile Leu Ile 210 215 220

Ala Lys Pro Ile Met Phe Arg Gly Glu Val Arg Leu Asn Val Glu Glu 235

Lys Lys Thr Arg Ala Ala Arg Glu Arg Glu Thr Arg Gly Gly Asp 250 245

Asp Arg Arg Asp Ile Arg Arg Asn Asp Arg Gly Pro Gly Pro Arg 265 260

Gly Ile Val Gly Gly Met Met Arg Asp Arg Asp Gly Arg Gly Pro 280 275

Pro Pro Arg Gly Gly Met Ala Gln Lys Leu Gly Ser Gly Arg Gly Thr 290

Gly Gln Met Glu Gly Arg Phe Thr Gly Gln Arg Arg 310 305

<210> 137

<211> 314

<212> PRT

<213> Homo sapien

<220>

<221> MISC\_FEATURE

<222> (4)..(4)

<223> X=any amino acid

<400> 137

Leu Gly Gly Xaa Glu Ser Gln Leu Leu Leu Ala Ser Gln Ser Glu Asp 10

Glu Val Glu Glu Glu Glu Glu Arg Gln Pro Ser Pro Glu Pro Val 25

Cln Glu Asn Ala Asn Ser Gly Tyr Tyr Glu Ala His Pro Val Thr Asn

Gly Ile Glu Glu Pro Leu Glu Glu Ser Ser His Glu Pro Glu Pro Glu

Pro Glu Ser Glu Thr Lys Thr Glu Glu Leu Lys Pro Gln Val Glu Glu 70 75

Lys Asn Leu Glu Glu Leu Glu Glu Lys Ser Thr Thr Pro Pro Pro Ala 90

Glu Pro Val Ser Leu Pro Gln Glu Pro Pro Lys Pro Arg Val Glu Ala 105

Lys Pro Glu Val Gln Ser Gln Pro Pro Arg Val Arg Glu Gln Arg Pro

Arg Glu Arg Pro Gly Phe Pro Pro Arg Gly Pro Arg Pro Gly Arg Gly 140

Asp Met Glu Gln Asn Asp Ser Asp Asn Arg Arg Ile Ile Arg Tyr Pro 150 155 145

Asp Ser His Gln Leu Phe Val Gly Asn Leu Pro His Asp Ile Asp Glu 170

Asn Glu Leu Lys Glu Phe Phe Met Ser Phe Gly Asn Val Val Glu Leu 185 180

Arg Ile Asn Thr Lys Gly Val Gly Gly Lys Leu Pro Asn Phe Gly Phe 200 195

Val Val Phe Asp Asp Ser Glu Pro Val Gln Arg Ile Leu Ile Ala Lys 215 210

Pro Ile Met Phe Arg Gly Glu Val Arg Leu Asn Val Glu Glu Lys Lys 230 225

Thr Arg Ala Ala Arg Glu Arg Glu Thr Arg Gly Gly Asp Asp Arg

Arg Asp Ile Arg Arg Asn Asp Arg Gly Pro Gly Gly Pro Arg Gly Ile 265 260

Val Gly Gly Met Met Arg Asp Arg Asp Gly Arg Gly Pro Pro Pro 275 280 285

Arg Gly Gly Met Ala Gln Lys Leu Gly Ser Gly Arg Gly Thr Gly Gln 300 290 295

Met Glu Gly Arg Phe Thr Gly Gln Arg Arg 310 305

<210> 138

<211> 169 <212> PRT

<213> Homo sapien

<400> 138

Met Glu Gln Asn Asp Ser Asp Asn Arg Arg Ile Ile Arg Tyr Pro Asp 10 1 5

Ser His Gln Leu Phe Val Gly Asn Leu Pro His Asp Ile Asp Glu Asn

Glu Leu Lys Glu Phe Phe Met Ser Phe Gly Asn Val Val Glu Leu Arg 40 45

Ile Asn Thr Lys Gly Val Gly Gly Lys Leu Pro Asn Phe Gly Phe Val 50 55

Val Phe Asp Asp Ser Glu Pro Val Gln Arg Ile Leu Ile Ala Lys Pro 70 75 80

Ile Met Phe Arg Gly Glu Val Arg Leu Asn Val Glu Glu Lys Lys Thr 85

Arg Ala Ala Arg Glu Arg Glu Thr Arg Gly Gly Asp Asp Arg Arg 105 100

Asp Ile Arg Arg Asn Asp Arg Gly Pro Gly Pro Arg Gly Ile Val 115 120

Gly Gly Met Met Arg Asp Arg Asp Gly Arg Gly Pro Pro Pro Arg 135 140 130

Gly Gly Met Ala Gln Lys Leu Gly Ser Gly Arg Gly Thr Gly Gln Met 145 150 155

Glu Gly Arg Phe Thr Gly Gln Arg Arg

<210> 139 <211> 147 <212> PRT <213> Homo sapien

<400> 139

Met Gly Arg Val Arg Thr Lys Thr Val Lys Lys Ala Ala Arg Val Ile 10

Ile Glu Lys Tyr Tyr Thr Arg Leu Gly Asn Asp Phe His Thr Asn Lys 25

Arg Val Cys Glu Glu Ile Ala Ile Ile Pro Ser Lys Lys Leu Arg Asn 35 40 45

Lys Ile Ala Gly Tyr Val Thr His Leu Met Lys Arg Ile Gln Arg Gly 50 55

Pro Val Arg Gly Ile Ser Ile Lys Leu Gln Glu Glu Glu Arg Glu Arg 65 70 75 80

Arg Asp Asn Tyr Val Pro Glu Val Ser Ala Leu Asp Gln Glu Ile Ile 85 90 95

Glu Val Asp Pro Asp Thr Lys Glu Met Leu Lys Leu Leu Asp Phe Gly
100 105 110

Ser Leu Ser Asn Leu Gln Val Ile His Pro Asn Cys Arg Leu Ser Asp 115 120 125

Leu Lys Val Gly Gln Thr Ala Val Gly Met Asn Phe Lys Thr Pro Arg 130 135 140

Gly Pro Val 145

<210> 140

<211> 166

<212> PRT

<213> Homo sapien

<220>

<221> MISC FEATURE

<222> (129)..(129)

<223> X=any amino acid

<220>

<221> MISC\_FEATURE

<222> (134)..(134)

<223> X=any amino acid

<400> 140

Ala Leu Thr Gly Phe Ala Cys Ala Ser Cys Phe Leu Phe Tyr Gln Gly
1 5 10 15

Pro Ala Asn Met Gly Arg Val Arg Thr Lys Thr Val Lys Lys Ala Ala 20 25 30

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199

Arg Val Ile Ile Glu Lys Tyr Tyr Thr Arg Leu Gly Asn Asp Phe His 40 \_45

Thr Asn Lys Arg Val Cys Glu Glu Ile Ala Ile Ile Pro Ser Lys Lys 55

Leu Arg Asn Lys Ile Ala Gly Tyr Val Thr His Leu Met Lys Arg Ile 75

Gln Arg Gly Pro Val Arg Gly Ile Ser Ile Lys Leu Gln Glu Glu 85 90

Arg Glu Arg Arg Asp Asn Tyr Val Pro Glu Val Ser Ala Leu Asp Gln 100 105

Glu Ile Ile Glu Val Asp Pro Asp Thr Lys Glu Met Leu Lys Leu Leu 115 120 125

Xaa Phe Gly Ser Leu Xaa Asn Leu Gln Val Ile His Pro Asn Cys Arg 130 135 140

Leu Ser Asp Leu Lys Val Gly Gln Thr Ala Val Gly Met Asn Phe Lys 145 150 155

Thr Pro Arg Gly Pro Val 165

<210> 141

<211> 254

<212> PRT <213> Homo sapien

<400> 141

Met Ser Val Asn Ala Ile Arg Lys Gln Ser Thr Asp Glu Glu Val Thr

Ser Leu Ala Lys Ser Leu Ile Lys Ser Trp Lys Lys Leu Leu Asp Gly 25

Pro Ser Thr Glu Lys Asp Leu Asp Glu Lys Lys Glu Pro Ala Ile 40

Thr Ser Gln Asn Ser Pro Glu Ala Arg Glu Glu Ser Thr Ser Ser Gly 55

Asn Val Ser Asn Arg Lys Asp Glu Thr Asn Ala Arg Asp Thr Tyr Val 70

Ser Ser Phe Pro Arg Ala Pro Ser Thr Ser Asp Ser Val Arg Leu Lys 85 90 95

Cys Arg Glu Met Leu Ala Ala Ala Leu Arg Thr Gly Asp Asp Tyr Ile 100 105 110

Ala Ile Gly Ala Asp Glu Glu Leu Gly Ser Gln Ile Glu Glu Ala 115 120 125

Ile Tyr Gln Glu Ile Arg Asn Thr Asp Met Lys Tyr Lys Asn Arg Val 130 , 135 140

Arg Ser Arg Ile Ser Asn Leu Lys Asp Ala Lys Asn Pro Asn Leu Arg 145 150 155 160

Lys Asn Val Leu Cys Gly Asn Ile Pro Pro Asp Leu Phe Ala Arg Met 165  $$170\$ 

Thr Ala Glu Glu Met Ala Ser Asp Glu Leu Lys Glu Met Arg Lys Asn 180 185 190

Leu Thr Lys Glu Ala Ile Arg Glu His Gln Met Ala Lys Thr Gly Gly
195 200 205

Thr Gln Thr Asp Leu Phe Thr Cys Gly Lys Cys Lys Lys Lys Asn Cys 210 215 220

Thr Tyr Thr Gln Val Gln Thr Arg Ser Ala Asp Glu Pro Met Thr Thr 225 230 235 240

Phe Val Val Cys Asn Glu Cys Gly Asn Arg Trp Lys Phe Cys 245 250

<210> 142

<211> 302

<212> PRT

<213> Homo sapien

<400> 142

Arg Gly Leu Asn Val Arg Leu Val Ile Ser Thr Val Leu His Val Cys 1 5 10 15

Leu Ala Ile Lys Asr Ala Ala Gly Ala Leu Asp Leu Lys Glu Leu 20 25 30

Lys Asn Ile Pro Met Thr Leu Glu Leu Gln Ser Thr Arg Ile Gly

Met Ser Val Asn Ala Ile Arg Lys Gln Ser Thr Asp Glu Glu Val Thr 50 55 60

Ser Leu Ala Lys Ser Leu Ile Lys Ser Trp Lys Lys Leu Leu Asp Gly 70 75 80

Pro Ser Thr Glu Lys Asp Leu Asp Glu Lys Lys Lys Glu Pro Ala Ile 85 90 95

Thr Ser Gln Asn Ser Pro Glu Ala Arg Glu Glu Ser Thr Ser Ser Gly 100 105 110

Asn Val Ser Asn Arg Lys Asp Glu Thr Asn Ala Arg Asp Thr Tyr Val 115 120 125

Ser Ser Phe Pro Arg Ala Pro Ser Thr Ser Asp Ser Val Arg Leu Lys 130 135 140

Cys Arg Glu Met Leu Ala Ala Leu Arg Thr Gly Asp Asp Tyr Ile 145 150 155 160

Ala Ile Gly Ala Asp Glu Glu Glu Leu Gly Ser Gln Ile Glu Glu Ala 165 170 175

Ile Tyr Gln Glu Ile Arg Asn Thr Asp Met Lys Tyr Lys Asn Arg Val

Arg Ser Arg Ile Ser Asn Leu Lys Asp Ala Lys Asn Pro Asn Leu Arg 195 200 205

Lys Asn Val Leu Cys Gly Asn Ile Pro Pro Asp Leu Phe Ala Arg Met 210 215 220

Thr Ala Glu Glu Met Ala Ser Asp Glu Leu Lys Glu Met Arg Lys Asn 225 230 235 240

Leu Thr Lys Glu Ala Ile Arg Glu His Gln Met Ala Lys Thr Gly Gly
245 250 255

Thr Gln Thr Asp Leu Phe Thr Cys Gly Lys Cys Lys Lys Lys Asn Cys 260 265 270

Thr Tyr Thr Gln Val Gln Thr Arg Ser Ala Asp Glu Pro Met Thr Thr

202

275 285 280

Phe Val Val Cys Asn Glu Cys Gly Asn Arg Trp Lys Phe Cys 290 295

<210> 143

<211> 225 <212> PRT

<213> Homo sapien

<400> 143

Met Val Ser His Ser Glu Leu Arg Lys Leu Phe Tyr Ser Ala Asp Ala

Val Cys Phe Asp Val Asp Ser Thr Val Ile Arg Glu Glu Gly Ile Asp 20 25 30

Glu Leu Ala Lys Ile Cys Gly Val Glu Asp Ala Val Ser Glu Met Thr 35 40 45

Arg Arg Ala Met Gly Gly Ala Val Pro Phe Lys Ala Ala Leu Thr Glu 50 55 60

Arg Leu Ala Leu Ile Gln Pro Ser Arg Glu Gln Val Gln Arg Leu Ile 70 75

Ala Glu Gln Pro Pro His Leu Thr Pro Gly Ile Arg Glu Leu Val Ser 85 90 95

Arg Leu Gln Glu Arg Asn Val Gln Val Phe Leu Ile Ser Gly Gly Phe 100 105 110

Arg Ser Ile Val Glu His Val Ala Ser Lys Leu Asn Ile Pro Ala Thr 115 120 125

Asn Val Phe Ala Asn Arg Leu Lys Phe Tyr Phe Asn Gly Glu Tyr Ala 130 135 140

Gly Phe Asp Glu Thr Gln Pro Thr Ala Glu Ser Gly Gly Lys Gly Lys 145 150 155 160

Val Ile Lys Leu Leu Lys Glu Lys Phe His Phe Lys Lys Ile Ile Met 165 170 175

Ile Gly Asp Gly Ala Thr Asp Met Glu Ala Cys Pro Pro Ala Asp Ala 180 185 190

Phe Ile Gly Phe Gly Gly Asn Val Ile Arg Gln Gln Val Lys Asp Asn 200

Ala Lys Trp Tyr Ile Thr Asp Phe Val Glu Leu Leu Gly Glu Leu Glu 215

Glu 225

<210> 144

<211> 249 <212> PRT

<213> Homo sapien

<400> 144

Met Lys Gln Thr Asn Ile Gln Lys Asn Thr Asn Thr Arg Asp Thr Ser 5 10

Lys Lys Thr Lys Asp Gln Leu Ile Ile Asp Ala Gly Gln Lys His Phe 25 30 20

Gly Ala Thr Val Cys Lys Ser Cys Gly Met Ile Tyr Thr Ala Ser Asn 40

Pro Glu Asp Glu Met Gln His Val Gln His His His Arg Phe Leu Glu 50

Gly Ile Lys Tyr Val Gly Trp Lys Lys Glu Arg Val Val Ala Glu Phe 70 75

Trp Asp Gly Lys Ile Val Leu Val Leu Pro His Asp Pro Ser Phe Ala

Ile Lys Lys Val Glu Asp Val Gln Glu Leu Val Asp Asn Glu Leu Gly 100 105

Phe Gln Gln Val Val Pro Lys Cys Pro Asn Lys Ile Lys Thr Phe Leu 115 120

Phe Ile Ser Asp Glu Lys Arg Val Val Gly Cys Leu Ile Ala Glu Pro 130 135

Ile Lys Gln Ala Phe Arg Val Leu Ser Glu Pro Ile Gly Pro Glu Ser 150 155

Pro Ser Ser Thr Glu Cys Pro Arg Ala Trp Gln Cys Ser Asp Val Pro

204

165 170 175

Glu Pro Ala Val Cys Gly Ile Ser Arg Ile Trp Val Phe Arg Leu Lys 180 185 190

Arg Arg Lys Arg Ile Ala Arg Arg Leu Val Asp Thr Leu Arg Asn Cys 195 200 205

Phe Met Phe Gly Cys Phe Leu Ser Thr Asp Glu Ile Ala Phe Ser Asp 210 215 220

Pro Thr Pro Asp Gly Lys Leu Phe Ala Thr Lys Tyr Cys Asn Thr Pro 225 230 235 235

Asn Phe Leu Val Tyr Asn Phe Asn Ser 245

<210> 145

<211> 113

<212> PRT

<213> Homo sapien

<400> 145

Met Lys Ser Phe Ser Lys Ser Ser Asn Lys Cys Thr Leu Asn Thr Ser 1 5 10 15

Thr Val Arg Glu Phe Leu Ser Phe Arg Met Asn Ala Ile His Thr Lys 20 25 30

Glu Leu Leu Thr Ser His Leu Gln Ser Pro Pro Gly His Arg Gln 35 40 45

Asp Pro Phe Asn Lys Ser Ser Ser Glu Thr Pro Ile Val Gln Asn Leu 50 55 60

Gln Leu Ala Thr Gly Tyr His His Ser Leu Trp Leu Cys Lys Ile Lys 65 70 75 80

Asp Leu Glu Glu Gly Trp Gly Gly Gly Ser Tyr Glu Lys Arg Gln Glu 85 90 95

Lys Ser Ser Phe Asp Pro Met Leu Ser Glu Ser Val His Glu Glu Glu 100 105 110

Ser

205

<210> 146 <211> 102 <212> PRT <213> Homo sapien

1220

<400> 146

Met Val Thr Glu Glu Lys Arg Ser Glu Ala Arg Glu Asn Glu Arg Ser 1 5 10 15

Leu Ala Phe Val Lys Met Val Gly His His Val Ala Phe Leu Glu Ala 20 25. 30

Asp Val Leu Gln Ala Glu Arg Asp His Gly Ala Phe Pro Gln Ala Leu 35 40 45

Arg Arg Trp Leu Gly Ser Ala Gly Leu Pro Ser Phe Arg Asn Lys Ser 50 55 60

Pro Ala Pro Val Pro Val Thr Tyr Glu Leu Pro Thr Leu Tyr Arg Thr 65 . 70 . 75 . 80

Glu Asp Tyr Phe Pro Val Asp Ala Gly Glu Ala Gln His His Pro Arg 85 90 95

Thr Cys Pro Arg Pro Leu 100

<210> 147

<211> 412

<212> PRT

<213> Homo sapien

<400> 147

Met Thr His Arg Arg Phe Lys Val Thr Ser Thr Val Ala Ala Ser 1 5 10 15

Leu Leu Pro Leu Gln Asp Glu Lys Glu Val Leu Leu Cys Lys Pro Ala 20 25 30

Trp Leu Ser Pro Ser Gly Thr Arg Thr Gly Gly Phe Leu Ala Val Pro 35 40 45

Gly Pro Pro Leu Arg Ala Lys Gly Pro Pro Val Leu Trp Pro Pro Pro 50 55 60

Ala His Pro Pro Arg Val Pro Gly Arg Glu His Ser Arg Trp Gly Arg 65 70 75 80

- Ser Pro Pro Ala Gln Arg Ala Ala Leu Gly Leu Arg Pro Tyr Leu Leu
- Leu Leu Pro Pro Ala Gln Leu Phe Asn Val Tyr Pro Trp Leu Gly 105 110
- Ala Leu Leu Gln Leu His Arg Pro Val Leu Arg Lys Ile Glu Glu Val 120
- Arg Ala Ile Leu Arg Thr Leu Leu Glu Ala Arg Arg Pro His Val Cys 135
- Pro Gly Asp Pro Val Cys Ser Tyr Val Asp Ala Leu Ile Gln Gln Gly 155 150
- Gln Gly Asp Asp Pro Glu Gly Leu Phe Ala Glu Ala Asn Ala Val Ala 170
- Cys Thr Leu Asp Met Val Met Ala Gly Thr Glu Thr Thr Ser Ala Thr 180 185
- Leu Gln Trp Ala Ala Leu Leu Met Gly Arg His Pro Asp Val Gln Gly 195 200
- Glu Thr Pro Ala Pro Gly Glu Thr Ala Pro Ser Ala Pro Gly Gly Pro 210 215
- Pro Gly Thr Arg Asp Gly Ala Ala Thr Gln Ala Ala Gln Pro Phe Ala 230
- Pro Gly Arg Val Gln Glu Glu Leu Asp Arg Val Leu Gly Pro Gly Arg 245 250 255
- Thr Pro Arg Leu Glu Asp Gln Gln Ala Leu Pro Tyr Thr Ser Ala Val 260 265 270
- Leu His Glu Val Gln Arg Phe Ile Thr Leu Leu Pro His Val Pro Arg 275 280
- Cys Thr Ala Ala Asp Thr Gln Leu Gly Gly Phe Leu Leu Pro Lys Gly 300 -290 295
- Thr Pro Val Ile Pro Leu Leu Thr Ser Val Leu Leu Asp Glu Thr Gln 315 310 305

Trp Gln Thr Pro Gly Gln Phe Asn Pro Gly His Phe Leu Asp Ala Asn 325 330 335

Gly His Phe Val Lys Arg Glu Ala Phe Leu Pro Phe Ser Ala Gly Arg 340 345 350

Arg Val Cys Val Gly Glu Arg Leu Ala Arg Thr Glu Leu Phe Leu Leu 355 360 365

Phe Ala Gly Leu Leu Gln Arg Tyr Arg Leu Leu Pro Pro Pro Gly Val 370 375 380

Ser Pro Ala Ser Leu Asp Thr Thr Pro Ala Arg Ala Phe Thr Met Arg 385 390 395 400

Pro Arg Ala Gln Ala Leu Cys Ala Val Pro Arg Pro 405 410

<210> 148

<211> 203

<212> PRT

<213> Homo sapien

<400> 148

Asp Pro Gly Ala Trp Arg Asp Gly Ser Phe Cys Pro Arg Gly Thr Pro 1 5 10 15

Arg Asp Glu Gly Trp Arg Cys His Pro Ser Gly Pro Pro Phe Ala Pro 20 25 30

Gly Arg Val Gln Glu Glu Leu Asp Arg Val Leu Gly Pro Gly Arg Thr 35 40 45

Pro Arg Leu Glu Asp Gln Gln Ala Leu Pro Tyr Thr Ser Ala Val Leu 50 55 60

His Glu Val Gln Arg Phe Ile Thr Leu Leu Pro His Val Pro Arg Cys 65 70 75 80

Thr Ala Ala Asp Thr Gln Leu Gly Gly Phe Leu Leu Pro Lys Gly Thr 85 90 95

Pro Val Ile Pro Leu Leu Thr Ser Val Leu Leu Asp Glu Thr Gln Trp 100 105 110

Gln Thr Pro Gly Gln Phe Asn Pro Gly His Phe Leu Asp Ala Asn Gly

208

115 120 125

His Phe Val Lys Arg Glu Ala Phe Leu Pro Phe Ser Ala Gly Arg Arg 130 135 140

Val Cys Val Gly Glu Arg Leu Ala Arg Thr Glu Leu Phe Leu Leu Phe 145 150 155

Ala Gly Leu Leu Gln Arg Tyr Arg Leu Leu Pro Pro Pro Gly Val Ser 165 170 175

Pro Ala Ser Leu Asp Thr Thr Pro Ala Arg Ala Phe Thr Met Arg Pro 180 185 190

Arg Ala Gln Ala Leu Cys Ala Val Pro Arg Pro 195 200

<210> 149

<211> 116

<212> PRT

<213> Homo sapien

<400> 149

Met Ala Arg Asp Ile Val Ala Met Ser Arg Ala Met Cys Leu Met Leu 1 5 10 15

Leu Ser Val Ala Arg Ala Phe Leu Leu Met Val Val Arg Thr Glu Glu 20 25 30

Val Ala Gly Phe Arg Trp Pro Asp Leu Arg Phe Asn Asp His His Asp 35 40 45

Thr Phe Ala Val Gly Cys Arg Leu His Ala His Ser Leu Ala Val Asn 50 60

Gln Ser Val Val Ala Glu Gly Ile Ala Gly Pro Gln Val Ile Gly Leu 65 70 75 80

Ser Ala Val Val Phe Gly Leu Ser Phe Glu Asn Met Glu Asn Trp Ser 85 90 95

Ser Ser Ala Arg Pro Ile Gln Leu Leu Met Pro Glu His Arg Tyr Ala 100 105 110

Asp Ile Arg Gln 115

<210> 150 <211> 141 <212> PRT

<213> Homo sapien

<400> 150

Gly Glu Arg Pro Leu Ser Trp Ser Pro Leu Gly Arg Gly His Leu Cys 10 5

Leu Val Pro Leu Gly Gly Arg Arg Gly Ala Cys Ala Gly Lys Ser Arg 25 30 2.0

Arg Pro Arg Trp Ala Asp His Glu Val Arg Ser Ser Arg Pro Ala Trp 40 45 35

Pro Thr Trp His Thr Trp His Ala His Arg Gly Asp Val Ala Cys His 50 55

Val Ser Asp Ala Ala Glu Arg Gly Ala Ser Ile Leu Val Asp Gly Gly 70 75 80

Pro His Ile Gly Gly Gly Arg Leu Pro Leu Ala Gly Ser Pro Leu Asn 90 95

Asp His His Asp Thr Phe Ala Val Gly Cys Arg Leu His Ala His Ser 100 105 110

Leu Ala Val Asn Gln Ser Val Val Ala Glu Gly Ile Ala Gly Pro Gln 115

Val Ile Gly Leu Ser Ala Val Val Phe Gly Leu Ser Phe 130 135

<210> 151

<211> 426 <212> PRT

<213> Homo sapien

<400> 151

Met Ser Pro Ala Pro Asp Ala Ala Pro Ala Pro Ala Ser Ile Ser Leu

Phe Asp Leu Ser Ala Asp Ala Pro Val Phe Gln Gly Leu Ser Leu Val

Ser His Ala Pro Gly Glu Ala Leu Ala Arg Ala Pro Arg Thr Ser Cys 40

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- Ser Gly Ser Gly Glu Arg Glu Ser Pro Glu Arg Lys Leu Leu Gln Gly 50 55 60
- Pro Met Asp Ile Ser Glu Lys Leu Phe Cys Ser Thr Cys Asp Gln Thr 65 70 75 80
- Phe Gln Asn His Gln Glu Gln Arg Glu His Tyr Lys Leu Asp Trp His 85 90 95
- Arg Phe Asn Leu Lys Gln Arg Leu Lys Asp Lys Pro Leu Leu Ser Ala 100 105 110
- Leu Asp Phe Glu Lys Gln Ser Ser Thr Gly Asp Leu Ser Ser Ile Ser 115 120 125
- Gly Ser Glu Asp Ser Asp Ser Ala Ser Glu Glu Asp Leu Gln Thr Leu 130 135 140
- Asp Arg Glu Arg Ala Thr Phe Glu Lys Leu Ser Arg Pro Pro Gly Phe 145 150 155 160
- Tyr Pro His Arg Val Leu Phe Gln Asn Ala Gln Gly Gln Phe Leu Tyr 165 170 175
- Ala Tyr Arg Cys Val Leu Gly Pro His Gln Asp Pro Pro Glu Glu Ala 180 185 190
- Glu Leu Leu Gln Asn Leu Gln Ser Arg Gly Pro Arg Asp Cys Val 195 200 205
- Val Leu Met Ala Ala Ala Gly His Phe Ala Gly Ala Ile Phe Gln Gly 210 215 220
- Arg Glu Val Val Thr His Lys Thr Phe His Arg Tyr Thr Val Arg Ala 225 230 235 240
- Lys Arg Gly Thr Ala Gln Gly Leu Arg Asp Ala Arg Gly Gly Pro Ser 245 250 255
- His Ser Ala Gly Ala Asn Leu Arg Arg Tyr Asn Glu Ala Thr Leu Tyr 260 265 270
- Lys Asp Val Arg Asp Leu Leu Ala Gly Pro Ser Trp Ala Lys Ala Leu 275 280 285

Glu Glu Ala Gly Thr Ile Leu Leu Arg Ala Pro Arg Ser Gly Arg Ser

Leu Phe Phe Gly Gly Lys Gly Ala Pro Leu Gln Arg Gly Asp Pro Arg 310

Leu Trp Asp Ile Pro Leu Ala Thr Arg Arg Pro Thr Phe Gln Glu Leu 325 330

Gln Arg Val Leu His Lys Leu Thr Thr Leu His Val Tyr Glu Glu Asp 345

Pro Arg Glu Ala Val Arg Leu His Ser Pro Gln Thr His Trp Lys Thr 360

Val Arg Glu Glu Arg Lys Lys Pro Thr Glu Glu Glu Ile Arg Lys Ile 375

Cys Arg Asp Glu Lys Glu Ala Leu Gly Gln Asn Glu Glu Ser Pro Lys 390 395

Gln Gly Leu Ile Thr Ile Trp Gln Leu Ser Asp Leu Ser Phe Cys Pro 410 405

Lys Asn Ala Leu Ala Asn Ser Leu Leu Ser 420

<210> 152 <211> 370 <212> PRT <213> Homo sapien

<400> 152

Met Ser Pro Ala Pro Asp Ala Ala Pro Ala Pro Ala Ser Ile Ser Leu

Phe Asp Leu Ser Ala Asp Ala Pro Val Phe Gln Gly Leu Ser Leu Arg

Glu His Tyr Lys Leu Asp Trp His Arg Phe Asn Leu Lys Gln Arg Leu 35 40

Lys Asp Lys Pro Leu Leu Ser Ala Leu Asp Phe Glu Lys Gln Ser Ser 50

Thr Gly Asp Leu Ser Ser Ile Ser Gly Ser Glu Asp Ser Asp Ser Ala

212 65 70 75 80 Ser Glu Glu Asp Leu Gln Thr Leu Asp Arg Glu Arg Ala Thr Phe Glu 90 Lys Leu Ser Arg Pro Pro Gly Phe Tyr Pro His Arg Val Leu Phe Gln Asn Ala Gln Gly Gln Phe Leu Tyr Ala Tyr Arg Cys Val Leu Gly Pro 115 His Gln Asp Pro Pro Glu Glu Ala Glu Leu Leu Gln Asn Leu Gln 135 Ser Arg Gly Pro Arg Asp Cys Val Val Leu Met Ala Ala Ala Gly His 150 155 Phe Ala Gly Ala Ile Phe Gln Gly Arg Glu Val Val Thr His Lys Thr 170 Phe His Arg Tyr Thr Val Arg Ala Lys Arg Gly Thr Ala Gln Gly Leu Arg Asp Ala Arg Gly Gly Pro Ser His Ser Ala Gly Ala Asn Leu Arg 200 Arg Tyr Asn Glu Ala Thr Leu Tyr Lys Asp Val Arg Asp Leu Leu Ala 215 Gly Pro Ser Trp Ala Lys Ala Leu Glu Glu Ala Gly Thr Ile Leu Leu 225 Arg Ala Pro Arg Ser Gly Arg Ser Leu Phe Phe Gly Gly Lys Gly Ala 245 Pro Leu Gln Arg Gly Asp Pro Arg Leu Trp Asp Ile Pro Leu Ala Thr 250 265 Arg Arg Pro Thr Phe Gln Glu Leu Gln Arg Val Leu His Lys Leu Thr 275 280 285 Thr Leu His Val Tyr Glu Glu Asp Pro Arg Glu Ala Val Arg Leu His 295 Ser Pro Gln Thr His Trp Lys Thr Val Arg Glu Glu Arg Lys Lys Pro 310

Thr Glu Glu Glu Ile Arg Lys Ile Cys Arg Asp Glu Lys Glu Ala Leu 325 330 335

Gly Gln Asn Glu Glu Ser Pro Lys Gln Gly Leu Ile Thr Ile Trp Gln 340 345 350

Leu Ser Asp Leu Ser Phe Cys Pro Lys Asn Ala Leu Ala Asn Ser Leu 355 360 365

Leu Ser 370

<210> 153

<211> 208

<212> PRT

<213> Homo sapien

<400> 153

Met Ser Pro Ala Pro Asp Ala Ala Pro Ala Pro Ala Ser Ile Ser Leu 1 5 10 15

Phe Asp Leu Ser Ala Asp Ala Pro Val Phe Gln Gly Leu Ser Leu Val 20 25 30

Ser His Ala Pro Gly Glu Ala Leu Ala Arg Ala Pro Arg Thr Ser Cys . 35 40 45

Ser Gly Ser Gly Glu Arg Glu Ser Pro Glu Arg Lys Leu Leu Gln Gly 50 55

Pro Met Asp Ile Ser Glu Lys Leu Phe Cys Ser Thr Cys Asp Gln Thr 65 70 75 80

Phe Gln Asn His Gln Glu Gln Arg Glu His Tyr Lys Leu Asp Trp His 85 90 95

Arg Phe Asn Leu Lys Gln Arg Leu Lys Asp Lys Pro Leu Leu Ser Ala 100 105 110

Leu Asp Phe Glu Lys Gln Ser Ser Thr Gly Asp Leu Ser Ser Ile Ser 115 120 125

Gly Ser Glu Asp Ser Asp Ser Ala Ser Glu Glu Asp Leu Gln Thr Leu 130 135 140

214

Asp Arg Glu Arg Ala Thr Phe Glu Lys Leu Ser Arg Pro Pro Gly Phe 145 150 150 160

Tyr Pro His Arg Val Leu Phe Gln Asn Ala Gln Gly Gln Phe Leu Tyr 165 170 175

Ala Tyr Arg Cys Val Leu Gly Pro His Gln Arg Gln Val Thr Val Gln 180 185 190

Val Ala Trp Leu Thr Pro Ala Phe Cys Thr Pro Ser Leu Asp Phe Pro 195 200 205

<210> 154

<211> 209

<212> PRT

<213> Homo sapien

<400> 154

Trp Thr Gln Leu Leu Met Cys Tyr Phe Tyr Leu Gly Asp Lys Ile Lys
1 10 15

Thr Ile Ser Phe Gln Ala Phe Ile Leu Met His Leu Leu Leu Pro Ser 20 25 30

Glu Tyr Ser Leu Asp Gly Phe His Met Ser Gly Phe Ser Leu Gly Ser 35 40 45

Gly Ser Glu Gly Glu Asp Gly Phe Gln Val Glu Leu Glu Leu Val Glu 50 55 60

Leu Thr Val Gly Thr Leu Asp Leu Cys Glu Ser Glu Val Leu Pro Lys 65 70 75 80

Arg Arg Arg Lys Arg Asn Lys Lys Glu Lys Ser Arg Asp Gln Glu 85 90 95

Ala Gly Ala His Arg Thr Leu Leu Gln Gln Thr Gln Glu Glu Glu Pro 100 105 110

Ser Thr Gln Ser Ser Gln Ala Val Ala Ala Pro Leu Gly Pro Leu Leu 115 120 125

Asp Glu Ala Lys Ala Pro Gly Gln Pro Glu Leu Trp Asn Ala Leu Leu 130 135 140

Ala Ala Cys Arg Ala Gly Asp Val Gly Val Leu Lys Leu Gln Leu Ala 145 150 155 160 Pro Ser Pro Ala Asp Pro Arg Val Leu Ser Leu Leu Ser Ala Pro Leu 165 170 175

Gly Ser Gly Gly Phe Thr Leu Leu His Ala Ala Ala Ala Ala Gly Arg 180 185 190

Gly Ser Val Val Arg Leu Leu Glu Ala Gly Ala Asp Pro Thr Val 195 200 205

Gln

<210> 155

<211> 125

<212> PRT

<213> Homo sapier

<400> 155

Met Ser Pro Ala Pro Asp Ala Ala Pro Ala Pro Ala Ser Ile Ser Leu 1 5 10 15

Phe Asp Leu Ser Ala Asp Ala Pro Val Phe Gln Gly Leu Ser Leu Val 20 25 30

Ser His Ala Pro Gly Glu Ala Leu Ala Arg Ala Pro Arg Thr Ser Cys  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Ser Gly Ser Glu Arg Glu Ser Pro Glu Arg Lys Leu Gln Gly 50 55 60

Pro Met Asp Ile Ser Glu Lys Leu Phe Cys Ser Thr Cys Asp Gln Thr 65 70 75 80

Phe Gln Asn His Gln Glu Gln Arg Glu His Tyr Lys Leu Asp Trp His

Arg Phe Asn Leu Lys Gln Arg Leu Lys Asp Lys Pro Leu Leu Ser Ala 100 105 110

Leu Asp Phe Glu Lys Gln Ser Ser Thr Gly Asp Glu Trp 115 120 125

<210> 156

<211> 191

<212> PRT

<213> Homo sapien

<400> 156

Glu Pro Ser Leu Asp Arg Pro Gly Asp Asp Gln Leu Val Leu Gly Gly
1 5 10 15

Gly Leu Cys Arg Val Glu Gly Ser Gln Val Pro Val Pro Ala Leu Ser 20 25 30

Pro Ala Thr Ala Pro Thr Ser Phe Glu Gly Pro Phe Gly Lys Ile Val 35 40

His Gln Val Arg Ala Ala Ile His Thr Pro Arg Phe Ser Lys Asp His 50 55 60

Lys Cys Ser Leu Val Phe Tyr Ile Leu Ser Pro His Phe Leu Asp Pro 65 70 75 80

Val Phe Leu Ser Thr Lys Ser His Ser Gln Arg Gln Pro Leu Leu Ala 85 90 95

Thr Leu Ser Ser Val Pro Gly Ala Pro Glu Pro Cys Pro Gln Asp Gly
100 105 110

Ser Pro Ala Ser His Pro Leu His Pro Pro Leu Cys Ile Ser Thr Gly
115 120 125

Ala Thr Val Pro Tyr Phe Ala Glu Gly Ser Gly Gly Pro Val Pro Thr 130 140

Thr Ser Thr Leu Ile Leu Pro Pro Glu Tyr Ser Ser Trp Gly Tyr Pro 145 150 155 160

Tyr Gly Glu Ser Thr Ala Arg Ala Trp Gln Gly Gly Asp Ala Lys Ser 165 170 175

Pro Thr Gln Thr Leu Leu Ser Ser Arg Arg Gly Pro Thr Val Leu 180 185 190

<210> 157

<211> 130

<212> PRT

<213> Homo sapien

<400> 157

Met Gly Cys Leu Leu Thr Gly Leu Pro Arg Thr Leu Pro Arg Trp Cys

1 10 15

Cys Leu Ala Pro Gly Arg Ile Pro Val Leu Ala Ala Ser Arg Gly Leu 20 25 30

Gly Arg Arg Leu Ala Gly Ala His Ala Ala Ile Pro Phe Ala Ala Ile 35 40 45

Arg Val Thr Cys Ile Gly Ser Cys Gly Val Ser Asn Lys Ala Asn Asp 50 55 60

Thr Ala Trp Val Val Glu Glu Gly Tyr Phe Asn Ser Ser Leu Ser Leu 65 70 75 80

Ala Asp Lys Gly Lys Phe Gly Ser Gln Phe Pro Ser Gly Asp Pro Trp 85 90 95

Gly Gln Pro Leu Glu Trp Gly Leu Ser Val Leu Ser Ser Pro Phe Pro 100 105 110

Arg Ser Ala Ser Trp Ile Trp His Trp Pro Ile Leu Ser Gln Gly Cys 115 120 125

Gly Pro 130

<210> 158

<211> 340

<212> PRT

<213> Homo sapien

<400> 158

Pro Gly Glu Ala His Phe Arg Glu Asp His Trp Pro Ala Ala Gly Pro 1 5 10 15

Thr Arg Arg Ser Ser Arg Pro Gly Val Pro Leu Gln Gly Ala Glu 20 . 25 30

Glu Asp Gly Ala Leu Trp Lys Gly Ala Arg Gly Phe Asn Gly Val Gln 35 40 45

Leu Phe Glu Gly Met Lys Ala Phe Lys Gly Lys Asp Gln Gln Val Arg 50 55 60

Leu Phe Arg Pro Trp Leu Asn Met Asp Arg Met Leu Arg Ser Ala Met 65 70 75 80

Arg Leu Cys Leu Pro Ser Phe Asp Lys Leu Glu Leu Leu Glu Cys Ile

95 90 85

Arg Arg Leu Ile Glu Val Asp Lys Asp Trp Val Pro Asp Ala Ala Gly 105

Thr Ser Leu Tyr Val Arg Pro Val Leu Ile Gly Asn Glu Pro Ser Leu 120

Gly Val Ser Gln Pro Thr Arg Ala Leu Leu Phe Val Ile Leu Cys Pro 135

Val Gly Ala Tyr Phe Pro Gly Gly Ser Val Thr Pro Val Ser Leu Leu 155

Ala Asp Pro Ala Phe Ile Arg Ala Trp Val Gly Gly Val Gly Asn Tyr 170

Lys Leu Gly Gly Asn Tyr Gly Pro Thr Val Leu Val Gln Gln Glu Ala 185 180

Leu Lys Arg Gly Cys Glu Gln Val Leu Trp Leu Tyr Gly Pro Asp His 200 195

Glm Leu Thr Glu Val Gly Thr Met Asm Ile Phe Val Tyr Trp Thr His 215 210

Glu Asp Gly Val Leu Glu Leu Val Thr Pro Pro Leu Asn Gly Val Ile 230 225

Leu Pro Gly Val Val Arg Gln Ser Leu Leu Asp Met Ala Gln Thr Trp

Gly Glu Phe Arg Val Val Glu Arg Thr Ile Thr Met Lys Gln Leu Leu 265

Arg Ala Leu Glu Glu Gly Arg Val Arg Glu Val Phe Gly Ser Gly Thr 280

Ala Cys Gln Val Cys Pro Val His Arg Ile Leu Tyr Lys Asp Arg Asn 295

Leu His Ile Pro Thr Met Glu Asn Gly Pro Glu Leu Ile Leu Arg Phe 310

Gln Lys Glu Leu Lys Glu Ile Gln Tyr Gly Ile Arg Ala His Glu Trp 330 325

Met Phe Pro Val 340

<210> 159

<211> 306

<212> PRT

<213> Homo sapien

<400> 159

Met Ala Pro Phe Gly Lys Glu His Glu Ala Leu Met Gly Glu Leu Phe 1 5 10 15

Glu Gly Met Lys Ala Phe Lys Gly Lys Asp Gln Gln Val Arg Leu Phe 20 25 30

Arg Pro Trp Leu Asn Met Asp Arg Met Leu Arg Ser Ala Met Arg Leu 35 40 45

Cys Leu Pro Ser Phe Asp Lys Leu Glu Leu Leu Glu Cys Ile Arg Arg 50 55 60

Leu Ile Glu Val Asp Lys Asp Trp Val Pro Asp Ala Ala Gly Thr Ser 65 70 75 80

Leu Tyr Val Arg Pro Val Leu Ile Gly Asn Glu Pro Ser Leu Gly Val 85 90 95

Ser Gln Pro Thr Arg Ala Leu Leu Phe Val Ile Leu Cys Pro Val Gly 100 105 110

Ala Tyr Phe Pro Gly Gly Ser Val Thr Pro Val Ser Leu Leu Ala Asp 115 120 125

Pro Ala Phe Ile Arg Ala Trp Val Gly Gly Val Gly Asn Tyr Lys Leu 130 135 140

Gly Gly Asn Tyr Gly Pro Thr Val Leu Val Gln Gln Glu Ala Leu Lys 145 150 155 160

Arg Gly Cys Glu Gln Val Leu Trp Leu Tyr Gly Pro Asp His Gln Leu 165 170 175

Thr Glu Val Gly Thr Met Asn Ile Phe Val Tyr Trp Thr His Glu Asp 180 185 190

220

Gly Val Leu Glu Leu Val Thr Pro Pro Leu Asn Gly Val Ile Leu Pro 195 200 205

Gly Val Val Arg Gln Ser Leu Leu Asp Met Ala Gln Thr Trp Gly Glu 210 215 220

Phe Arg Val Val Glu Arg Thr Ile Thr Met Lys Gln Leu Leu Arg Ala 225 230 235 240

Leu Glu Glu Gly Arg Val Arg Glu Val Phe Gly Ser Gly Thr Ala Cys 245 250 255

Gln Val Cys Pro Val His Arg Ile Leu Tyr Lys Asp Arg Asn Leu His 260 265 270

Ile Pro Thr Met Glu Asn Gly Pro Glu Leu Ile Leu Arg Phe Gln Lys 275 280 285

Glu Leu Lys Glu Ile Gln Tyr Gly Ile Arg Ala His Glu Trp Met Phe 290 295 300

Pro Val

<210> 160

<211> 485

<212> PRT

<213> Homo sapien

<400> 160

Gln Ile Val Tyr Leu Tyr Ile Gln Arg Ile Ile Arg Val Phe His Gly
1 5 10 15

Val Asn Ala Pro Asp Asn Pro Leu Lys Glu Glu His Leu Val Gln Leu 20 25 30

Asn Glu Thr Asp Ile Leu Arg Val Leu Asp Gly Asn Thr Gly Gly Thr 35 40 45

Tyr Gly Gly His Ile Pro Gly Ser Asp Arg Ala Gly Leu Asn Arg His 50 55 60

Asp Lys Ser Glu Asn Pro Gly Arg Ile Tyr Ala Gly Gly Ser Ser Thr 65 70 75 80

Ala Ala Gly Asp Pro Ser Leu Leu Ser Ala Ala Arg Ile Met Ala Ala 85 90 95

221

Ala Ala Leu Gly Gln Ile Trp Ala Arg Lys Leu Leu Ser Val Pro Trp 100 105 110

Leu Leu Cys Gly Pro Arg Arg Tyr Ala Ser Ser Ser Phe Lys Ala Ala 115 120 125

Asp Leu Gln Leu Glu Met Thr Gln Lys Pro His Lys Lys Pro Gly Pro 130 135 140

Gly Glu Pro Leu Val Phe Gly Lys Thr Phe Thr Asp His Met Leu Met 145 150 155 160

Val Glu Trp Asn Asp Lys Gly Trp Gly Gln Pro Arg Ile Gln Pro Phe 165 170 175

Gln Asn Leu Thr Leu His Pro Ala Ser Ser Ser Leu His Tyr Ser Leu 180 185 190

Gln Leu Phe Glu Gly Met Lys Ala Phe Lys Gly Lys Asp Gln Gln Val 195 200 205

Arg Leu Phe Arg Pro Trp Leu Asn Met Asp Arg Met Leu Arg Ser Ala 210 220

Met Arg Leu Cys Leu Pro Ser Phe Asp Lys Leu Glu Leu Leu Glu Cys 225 230 235

Ile Arg Arg Leu Ile Glu Val Asp Lys Asp Trp Val Pro Asp Ala Ala 245 250 255

Gly Thr Ser Leu Tyr Val Arg Pro Val Leu Ile Gly Asn Glu Pro Ser 260 265 270

Leu Gly Val Ser Gln Pro Thr Arg Ala Leu Leu Phe Val Ile Leu Cys 275 280 285

Pro Val Gly Ala Tyr Phe Pro Gly Gly Ser Val Thr Pro Val Ser Leu 290 295 300

Leu Ala Asp Pro Ala Phe Ile Arg Ala Trp Val Gly Gly Val Gly Asn 305 310 315

Tyr Lys Leu Gly Gly Asn Tyr Gly Pro Thr Val Leu Val Gln Glu 325 330 335

Ala Leu Lys Arg Gly Cys Glu Gln Val Leu Trp Leu Tyr Gly Pro Asp 340 345 350

His Gln Leu Thr Glu Val Gly Thr Met Asn Ile Phe Val Tyr Trp Thr 355 360 365

His Glu Asp Gly Val Leu Glu Leu Val Thr Pro Pro Leu Asn Gly Val

Ile Leu Pro Gly Val Val Arg Gln Ser Leu Leu Asp Met Ala Gln Thr 385 390 395 400

Trp Val Arg Thr Trp His Leu Leu Val Met Gly Ala Met Cys Gln Gly 405 410 415

Pro Gly His Gln Arg Ala Gly Thr Gly Ala His Trp His Val Ser Ala 420 425 430

Pro Ser Pro Gly Ser Val Ser Pro Val Gly Pro Leu Ser Phe Ser Leu 435 440 445

Ser Ser Gly Arg Glu Arg Trp Arg Ser Ala Ala Gly Gln Pro Ser Gly 450 455

Asp Thr Cys Leu Cys Gln Leu Pro Cys Arg Val Ser Ser Gly Trp Trp 465 470 475 480

Ser Ala Arg Ser Pro

<210> 161

<211> 465

<212> PRT

<213> Homo sapien

<400> 161

Met Ala Ala Ala Leu Gly Gln Ile Trp Ala Arg Lys Leu Leu Ser 1 5 10 15

Val Pro Trp Leu Leu Cys Gly Pro Arg Arg Tyr Ala Ser Ser Ser Phe 20 25 30

Lys Ala Ala Asp Leu Gln Leu Glu Met Thr Gln Lys Pro His Lys Lys 35 40 45

Pro Gly Pro Gly Glu Pro Leu Val Phe Gly Lys Thr Phe Thr Asp His

60 55 50 Met Leu Met Val Glu Trp Asn Asp Lys Gly Trp Gly Gln Pro Arg Ile 70 Gln Pro Phe Gln Asn Leu Thr Leu His Pro Ala Ser Ser Ser Leu His 90 Tyr Ser Leu Gln Leu Phe Glu Gly Met Lys Ala Phe Lys Gly Lys Asp . 105 Gln Gln Val Arg Leu Phe Arg Pro Trp Leu Asn Met Asp Arg Met Leu 120 Arg Ser Ala Met Arg Leu Cys Leu Pro Ser Phe Asp Lys Leu Glu Leu 135 Leu Glu Cys Ile Arg Arg Leu Ile Glu Val Asp Lys Asp Trp Val Pro 150 145 Asp Ala Ala Gly Thr Ser Leu Tyr Val Arg Pro Val Leu Ile Gly Asn 165 170 Glu Pro Ser Leu Gly Val Ser Gln Pro Thr Arg Ala Leu Leu Phe Val Ile Leu Cys Pro Val Gly Ala Tyr Phe Pro Gly Gly Ser Val Thr Pro

200

Val Ser Leu Leu Ala Asp Pro Ala Phe Ile Arg Ala Trp Val Gly Gly

Val Gly Asn Tyr Lys Leu Gly Gly Asn Tyr Gly Pro Thr Val Leu Val 235

Gln Gln Glu Ala Leu Lys Arg Gly Cys Glu Gln Val Leu Trp Leu Tyr 250 245

Gly Pro Asp His Gln Leu Thr Glu Val Gly Thr Met Asn Ile Phe Val 265 260

Tyr Trp Thr His Glu Asp Gly Val Leu Glu Leu Val Thr Pro Pro Leu 280 275

Asn Gly Val Ile Leu Pro Gly Val Val Arg Gln Ser Leu Leu Asp Met 295

224

Ala Gln Thr Trp Val Glu Asp Met Ala Ser Ser Gly Asp Gly Arg His 305

Val Pro Gly Ala Arg Ala Ser Glu Gly Trp Asp Trp Gly Thr Leu Ala 330

Cys Leu Cys Pro Phe Ser Trp Val Cys Leu Ser Arg Trp Ala Ser Val 345

Phe Leu Thr Ile Leu Arg Glu Gly Glu Val Glu Val Cys Ser Arg Ala 360 355

Thr Leu Trp Gly His Val Ser Leu Pro Thr Ala Leu Gln Gly Glu Phe 375

Arg Val Val Glu Arg Thr Ile Thr Met Lys Gln Leu Leu Arg Ala Leu 390 395

Glu Glu Gly Arg Val Arg Glu Val Phe Gly Ser Gly Thr Ala Cys Gln 405 410

Val Cys Pro Val His Arg Ile Leu Tyr Lys Asp Arg Asn Leu His Ile 425 420

Pro Thr Met Glu Asn Gly Pro Glu Leu Ile Leu Arg Phe Gln Lys Glu 440

Leu Lys Glu Ile Gln Tyr Gly Ile Arg Ala His Glu Trp Met Phe Pro 455

Val

465

<210> 162

<211> 74

<212> PRT

<213> Homo sapien

<220>

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<222> (3)..(5)

<223> X=any amino acid

<220>

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<222> (8)..(10) <223> X=any amino acid

<220>

<221> MISC\_FEATURE
<222> (13)..(14)
<223> X=any amino acid

<400> 162

Thr Leu Xaa Xaa Yaa Phe Asn Xaa Xaa Xaa Ser Ser Xaa Xaa Lys Ile 15 10 1 5

Arg Lys Asn Thr Ala Ser Tyr Val Pro Lys Glu Lys Lys Ile Lys Gly 20 25

Thr Met Pro Thr Cys Ser Thr Ile Lys Ala Ser Phe Ser Tyr Phe Phe 40 35

Asn Thr Lys Tyr Lys Gln Arg Ile His Ile Leu Lys Thr Glu Leu Arg 50 60

Ser Arg His Ala Val Leu Glu Thr Leu Gln 65 70

<210> 153

<211> 63

<212> PRT

<213> Homo sapien

<400> 163

Thr Arg Val Leu Gln Cys Ala Pro Arg Cys Ser Pro Asn Cys Val Ala 10

Phe Thr Ala Ala Val Thr Val Pro Ala Cys Ile Tyr Ala Leu Phe Gly 20 25

Pro Cys Glu His Thr Gly Ile Leu Val Ile Leu Pro Pro Met Glu Tyr 40

Leu Trp Arg Ser Pro Val Phe Ile Tyr Phe Gly Ile Asn Pro Leu

<210> 164

<211> 313

<212> PRT

<213> Homo sapien

<400> 164

Met Lys Cys Glu His Cys Thr Arg Lys Glu Cys Ser Lys Lys Thr Lys

226

1 5 10 15

Thr Asp Asp Gln Glu Asn Val Ser Ala Asp Ala Pro Ser Pro Ala Gln 20 25 30

Glu Asn Gly Glu Lys Gly Glu Phe His Lys Leu Ala Asp Ala Lys Ile 35 40 45

Phe Leu Ser Asp Cys Leu Ala Cys Asp Ser Cys Met Thr Ala Glu Glu 50 55 60

Gly Val Gln Leu Ser Gln Gln Asn Ala Lys Asp Phe Phe Arg Val Leu 65 70 75 80

Asn Leu Asn Lys Lys Cys Asp Thr Ser Lys His Lys Val Leu Val Val 85 90 95

Ser Val Cys Pro Gln Ser Leu Pro Tyr Phe Ala Ala Lys Phe Asn Leu 100 105 110

Ser Val Thr Asp Ala Ser Arg Arg Leu Cys Gly Phe Leu Lys Ser Leu 115 120 125

Gly Val His Tyr Val Phe Asp Thr Thr Ile Ala Ala Asp Phe Ser Ile 130 135 140

Glu Glu Arg Thr Leu Pro Met Leu Thr Ser Ala Cys Pro Gly Trp Val 165  $\,$  170  $\,$  175

Arg Tyr Ala Glu Arg Val Leu Gly Arg Pro Ile Thr Ala His Leu Cys 180 185 190

Thr Ala Lys Ser Pro Gln Gln Val Met Gly Ser Leu Val Lys Asp Tyr 195 200 205

Phe Ala Arg Gln Gln Asn Leu Ser Pro Glu Lys Ile Phe His Val Ile 210 215 220

Val Ala Pro Cys Tyr Asp Lys Lys Leu Glu Ala Leu Gln Glu Ser Leu 225 230 235 240

Pro Pro Ala Leu His Gly Ser Arg Gly Ala Asp Cys Val Leu Thr Ser 245 250 255

Gly Glu Ile Ala Gln Ile Met Glu Gln Gly Asp Leu Ser Val Arg Asp 260 265 270

Ala Ala Val Asp Thr Leu Val Ser Gly Phe Ser Gly Glu Ser Pro Leu 275 280 285

Gly Gly Arg Thr Ser Arg Gln Pro Cys Gln Pro Pro Ala Arg Pro Arg 290 295 300

Ala Ala Leu Leu Tyr Asp Glu Ala Met 305 310

<210> 165

<211> 395

<212> PRT

<213> Homo sapien

<400> 165

Glu Pro Arg Val Arg Arg Val Ser Asn Ala Glu Leu Ala Asp Arg Ala 1 5 10 15

Arg Pro Arg Pro Pro Arg Ala Gln Gly Pro Pro Gly Pro Val Thr Thr 20 25 30

Gly Pro Ser Thr Leu Glu Arg Pro Gln Leu Gly Leu Gly Thr Val Arg

Ala Leu Thr Asp Ser Leu Val Asn Ala Ala Trp Pro Pro Pro Pro Pro 50 55 60

Gln Asp Pro Arg Glu Ala Glu Thr Gly Ala Arg Thr Arg Ser Pro Arg 65 70 75 80

Arg Arg Thr Trp Ser Glu Pro Ala Ala Pro Pro Arg Ala Leu Arg Leu 85 90 95

Ala Leu Gly Pro Gly Pro Pro Leu Pro Asp Thr Val Ile Gly Leu Gly 100 105 110

Lys Ala Val Arg Val Gly Asn Pro Ile Gly Pro Gly Val Arg Leu Val

Leu Ser Arg Cys Ser His Trp Pro Ser Ala Ala Val Gly Glu Ala 130 135 140

228

Ala Ser Gly Glu Asp Asn Lys Gly Pro Arg Ala Ala Gly Ser Gly Val 145 150 155 160

Pro Val Ser Arg Cys Phe Pro Glu Ala Glu Ala Pro Gly Leu Pro Pro 165 170 175

Ala Ala Leu Gln Met Lys Cys Glu His Cys Thr Arg Lys Glu Cys Ser 180 185 190

Lys Lys Thr Lys Thr Asp Asp Gln Glu Asn Val Ser Ala Asp Ala Pro 195 200 205

Ser Pro Ala Gln Glu Asn Gly Glu Lys Gly Glu Phe His Lys Leu Ala 210 215 220

Asp Ala Lys Ile Phe Leu Ser Asp Cys Leu Ala Cys Asp Ser Cys Met 225 230 230 240

Thr Ala Glu Glu Gly Val Gln Leu Ser Gln Gln Asn Ala Lys Asp Phe 245 250 255

Phe Arg Val Leu Asn Leu Asn Lys Lys Cys Asp Thr Ser Lys His Lys 260 265 270

Val Leu Val Val Ser Val Cys Pro Gln Ser Leu Pro Tyr Phe Ala Ala 275 280 285

Lys Phe Asn Leu Ser Val Thr Asp Ala Ser Arg Arg Leu Cys Gly Phe 290 295 300

Leu Lys Ser Leu Gly Val His Tyr Val Phe Asp Thr Thr Ile Ala Ala 305 310 315 320

Asp Phe Ser Ile Leu Glu Ser Gln Lys Glu Phe Val Arg Arg Tyr Arg 325 330 335

Gln His Ser Glu Glu Glu Arg Thr Leu Pro Met Leu Thr Ser Ala Cys 340 345 350

Pro Gly Trp Val Arg Tyr Ala Glu Arg Val Leu Gly Arg Pro Ile Thr 355 360 365

Ala His Leu Cys Thr Ala Lys Ser Pro Gln Gln Val Met Gly Ser Leu 370 380

Val Lys Asp Tyr Phe Ala Arg Gln Gln Val Ser

229

395 390 385

<210> 166 <211> 285 <212> PRT <213> Homo sapien

<400> 166

Met Gly Ser Ala Phe Pro Val Leu Pro Gly Gly Ser Thr Gly Trp Gly 1 5 10 15

Ala Leu Gln Met Phe Gly Arg Thr Thr Pro Ser Pro Glu Gly Gly Ser

Arg Gln Thr Trp Met Glu Cys Trp Cys Ser Asn Leu Ser Pro Glu Lys 40

Ile Phe His Val Ile Val Ala Pro Cys Tyr Asp Lys Lys Leu Glu Ala 50 55 60

Leu Gln Glu Ser Leu Pro Pro Ala Leu His Gly Ser Arg Gly Ala Asp 75

Cys Val Leu Thr Ser Gly Glu Ile Ala Gln Ile Met Glu Gln Gly Asp 85 90 95

Leu Ser Val Arg Asp Ala Ala Val Asp Thr Leu Phe Gly Asp Leu Lys 100 105 110

Glu Asp Lys Val Thr Arg His Asp Gly Ala Ser Ser Asp Gly His Leu 120 125 115

Ala His Ile Phe Arg His Ala Ala Lys Glu Leu Phe Asn Glu Asp Val 130 135 140

Glu Glu Val Thr Tyr Arg Ala Leu Arg Asn Lys Asp Phe Gln Glu Val 145 150

Thr Leu Glu Lys Asn Gly Glu Val Val Leu Arg Phe Ala Ala Ala Tyr

Gly Phe Arg Asn Ile Gln Asn Met Ile Leu Lys Leu Lys Gly Lys 180 185 190

Phe Pro Phe His Phe Val Glu Val Leu Ala Cys Ala Gly Gly Cys Leu 195 200

Asn Gly Arg Gly Gln Ala Gln Thr Pro Asp Gly His Ala Asp Lys Ala 210 215 220

Leu Leu Arg Gln Met Glu Gly Ile Tyr Ala Asp Ile Pro Val Arg Arg 225 230 235 240

Pro Glu Ser Ser Ala His Val Gln Glu Leu Tyr Gln Glu Trp Leu Glu 245 250 255

Gly Ile Asn Ser Pro Lys Ala Arg Glu Val Leu His Thr Tyr Gln 260 265 270

Ser Gln Glu Arg Gly Thr His Ser Leu Asp Ile Lys Trp 275 280 285

<210> 167

<211> 170

<212> PRT

<213> Homo sapien

<400> 167

Asp Ser Val Ser His Pro Ala Lys Lys Phe Ala Met Ser Ala Ala Lys

1 10 15

Cys Lys Gln Ser Arg Leu Trp Val Lys Ala Leu Val Met Lys Asn Lys 20 25 30

Lys Lys Ile Pro Lys His Phe Leu Thr Leu Val Leu Lys Lys Thr 35 40 45

Pro Ser Pro Lys Val Cys Phe Ser Phe Asn Phe Leu Asp Phe Ala Arg 50 55 60

Leu Ser Gly Cys Lys Leu Pro Thr Leu Phe Trp Met Asp Arg Asp Lys 65 70 75 80

Val Phe Lys Gln Arg Leu Cys Pro Leu His Lys Pro Phe Pro Pro Pro 85 90 95

Pro Pro Gln Pro Pro Ala Ala Ile Ile Thr Gly Ala Val Lys Trp Leu 100 105 110

Leu Ser Asp Gly His Thr Thr Arg Arg Gln Met Lys Arg Ser Gly Ser 115 120 125

Lys Arg Gly Ser Ala Ser Gln Tyr Gln Pro Ala Val Pro Arg Gly Gly

231

130 135 140

Ser Ala Gly Arg Thr Val Phe Pro Arg Gln Ala Ala Met Pro Pro Pro 145 150 155 160

Thr Ala Lys Ala Pro Lys Ala Thr Ser Val 165 170

<210> 168

<211> 159

<212> PRT

<213> Homo sapien

<400> 168

Met Ser Ala Ala Lys Cys Lys Gln Ser Arg Leu Trp Val Lys Ala Leu 1 5 10 15

Val Met Lys Asn Lys Lys Lys Ile Pro Lys His Phe Leu Thr Leu 20 25 30

Val Leu Lys Lys Thr Pro Ser Pro Lys Val Cys Phe Ser Phe Asn Phe 35 40 45

Leu Asp Phe Ala Arg Leu Ser Gly Cys Lys Leu Pro Thr Leu Phe Trp 50 55 60

Met Asp Arg Asp Lys Val Phe Lys Gln Arg Leu Cys Pro Leu His Lys 65 70 75 80

Pro Phe Pro Pro Pro Pro Gln Pro Pro Ala Ala Ile Ile Thr Gly 85 90 95

Ala Val Lys Trp Leu Leu Ser Asp Gly His Thr Thr Arg Arg Gln Met 100  $$105\$ 

Lys Arg Ser Gly Ser Lys Arg Gly Ser Ala Ser Gln Tyr Gln Pro Ala 115 120 125

Val Pro Arg Gly Gly Ser Ala Gly Arg Thr Val Phe Pro Arg Gln Ala 130 135 140

Ala Met Pro Pro Pro Thr Ala Lys Ala Pro Lys Gly Asn Ile Arg 145 150 155

<210> 169

<211> 170

<212> PRT

<213> Homo sapien

<400> 169

Asp Ser Val Ser His Pro Ala Lys Lys Phe Ala Met Ser Ala Ala Lys 1 5 10

Cys Lys Gln Ser Arg Leu Trp Val Lys Ala Leu Val Met Lys Asn Lys 20 25 30

Lys Lys Lys Ile Pro Lys His Phe Leu Thr Leu Val Leu Lys Lys Thr 35 40

Pro Ser Pro Lys Val Cys Phe Ser Phe Asn Phe Leu Asp Phe Ala Arg 50 55

Leu Ser Gly Cys Lys Leu Pro Thr Leu Phe Trp Met Asp Arg Asp Lys 70 75 80

Val Phe Lys Gln Arg Leu Cys Pro Leu His Lys Pro Phe Pro Pro Pro 85 90 95

Pro Pro Gln Pro Pro Ala Ala Ile Ile Thr Gly Ala Val Lys Trp Leu 100 105 110

Leu Ser Asp Gly His Thr Thr Arg Arg Gln Met Lys Arg Ser Gly Ser

Lys Arg Gly Ser Ala Ser Gln Tyr Gln Pro Ala Val Pro Arg Gly Gly

Ser Ala Gly Arg Thr Val Phe Pro Arg Gln Ala Ala Met Pro Pro Pro 145 150 155 160

Thr Ala Lys Ala Pro Lys Ala Thr Ser Val

<210> 170 <211> 255 <212> PRT <213> Homo sapien

<400> 170

Gln Leu Leu Arg Asp Pro Asn Val Ala Leu Glu Leu Ser Ala Met Cys 10

Ser Thr Val Pro Trp Arg Arg Thr Leu Arg Glu Gly Gln Pro Cys His 25

Leu Ser Leu Pro His His His Ser Pro Pro Pro Ile Lys Leu Gln Ser 35 40 45

Gly Cys Trp Thr Pro Leu Gly Ala Val Ser Ala His His Pro Leu Cys 50 55

Ala Ala Thr Trp Ser Gln Ala His Cys Pro Leu Ala Gly Arg Gly Pro 75

Ser Arg Arg Arg Cys Gly Leu His Arg Ala Pro Ser Thr Lys Glu Ser 90 95

Ala Asn Ala Ser Ala Gly Pro Arg Ala Met Ala Ser Leu Pro Gln Leu 100 105 110

Met Ala Ala Pro Thr Ser Ser Cys Thr Ser Leu Met Trp Lys Gly Ser 115 120 125

Met Ser Gln Trp Lys Ala Tnr Arg Ser Pro Ile Lys Cys Ala Pro Ser 135 140

His Pro Arg Met Arg Ser Cys Arg Pro Trp Arg Ser Ser Ser Leu Thr 150 155 160 145

Trp His Gln Ala Pro Ser Met Arg Pro Gly Leu Asp Met Ser Ser Ala 165 170 175

Pro Arg Arg Trp Trp Lys His Pro Leu Ser Cys Ala Cys Gly Arg Leu 190 180 185

Cys Gly Glu Glu Ala Ala Asp Thr Gly Asp Asp Ile Leu Pro His Glu 195 200 205

Thr Gly Leu Gln Pro Gly Met Val Pro Leu Lys Tyr Leu Leu Glu Glu 210 215 220

Gly Val Trp Gly Ala Gly Val Gly Cys Gly Val Phe Pro Ala Ile Ser 240 230 235 225

Thr Ala Tyr Asp His Cys Asn Asn Leu Ser Pro Ser Glu Glu His 245 - 250

<210> 171 <211> 147 <212> PRT

<213> Homo sapien

<400> 171

Met Ser Ser Glu Pro Pro Pro Pro Pro Gln Pro Pro Thr His Gln Ala 1 5 10 15

Ser Val Gly Leu Leu Asp Thr Pro Arg Ser Arg Glu Arg Ser Pro Ser 20 25 30

Pro Leu Arg Gly Asn Val Val Pro Ser Pro Leu Pro Thr Arg Arg Thr 35 40 45

Arg Thr Phe Ser Ala Thr Val Arg Ala Ser Gln Gly Pro Val Tyr Lys 50 55 60

Gly Val Cys Lys Cys Phe Cys Arg Ser Lys Gly His Gly Phe Ile Thr 65 70 75 80

Pro Ala Asp Gly Gly Pro Asp Ile Phe Leu His Ile Ser Asp Val Glu 85 90 95

Gly Glu Tyr Val Pro Val Glu Gly Asp Glu Val Thr Tyr Lys Met Cys 100 105 110

Ser Ile Pro Pro Lys Asn Glu Lys Leu Gln Ala Val Glu Val Val Ile 115 120 125

Thr His Leu Ala Pro Gly Thr Lys His Glu Thr Trp Ser Gly His Val 130 135 140

Ile Ser Ser 145

<210> 172

<211> 255

<212> PRT

<213> Homo sapien

<400> 172

Gln Leu Leu Arg Asp Pro Asn Val Ala Leu Glu Leu Ser Ala Met Cys 1 5 10 15

Ser Thr Val Pro Trp Arg Arg Thr Leu Arg Glu Gly Gln Pro Cys His 20 25 30

Leu Ser Leu Pro His His His Ser Pro Pro Pro Ile Lys Leu Gln Ser 35 40 45 Gly Cys Trp Thr Pro Leu Gly Ala Val Ser Ala His His Pro Leu Cys 55 50

Ala Ala Thr Trp Ser Gln Ala His Cys Pro Leu Ala Gly Arg Gly Pro

Ser Arg Arg Arg Cys Gly Leu His Arg Ala Pro Ser Thr Lys Glu Ser 85

Ala Asn Ala Ser Ala Gly Pro Arg Ala Met Ala Ser Leu Pro Gln Leu 105

Met Ala Ala Pro Thr Ser Ser Cys Thr Ser Leu Met Trp Lys Gly Ser 115 120 125

Met Ser Gln Trp Lys Ala Thr Arg Ser Pro Ile Lys Cys Ala Pro Ser 130 135

His Pro Arg Met Arg Ser Cys Arg Pro Trp Arg Ser Ser Ser Leu Thr 145

Trp His Gln Ala Pro Ser Met Arg Pro Gly Leu Asp Met Ser Ser Ala 165 170

Pro Arg Arg Trp Trp Lys His Pro Leu Ser Cys Ala Cys Gly Arg Leu 185 180

Cys Gly Glu Glu Ala Ala Asp Thr Gly Asp Asp Ile Leu Pro His Glu 200 195

Thr Gly Leu Gln Pro Gly Met Val Pro Leu Lys Tyr Leu Leu Glu Glu 210

Gly Val Trp Gly Ala Gly Val Gly Cys Gly Val Phe Pro Ala Ile Ser 225 230

Thr Ala Tyr Asp His Cys Asn Asn Leu Ser Pro Ser Glu Glu His 250 245

<210> 173

<211> 243 <212> PRT <213> Homo sapien

<400> 173

Leu Arg Thr Gly Arg Asn Ser Gly Gly Gly Gln Asn Gly Leu Gln Gly
1 5 10 15

Gln Pro Cys His Leu Ser Leu Pro His His Ser Pro Pro Pro Ile 20 25 30

Lys Leu Gln Ser Gly Cys Trp Thr Pro Leu Gly Ala Val Ser Ala His 35 40 45

His Pro Leu Cys Ala Ala Thr Trp Ser Gln Ala His Cys Pro Leu Ala 50 55 60

Gly Arg Gly Pro Ser Arg Arg Cys Gly Leu His Arg Ala Pro Ser 65 70 75 80

Thr Lys Glu Ser Ala Asn Ala Ser Ala Gly Pro Arg Ala Met Ala Ser 85 90 95

Leu Pro Gln Leu Met Ala Ala Pro Thr Ser Ser Cys Thr Ser Leu Met 100 105 110

Trp Lys Gly Ser Met Ser Gln Trp Lys Ala Thr Arg Ser Pro Ile Lys 115 120 125

Cys Ala Pro Ser His Pro Arg Met Arg Ser Cys Arg Pro Trp Arg Ser 130 135 140

Ser Ser Leu Thr Trp His Gln Ala Pro Ser Met Arg Pro Gly Leu Asp 145 150 155 160

Met Ser Ser Ala Pro Arg Arg Trp Trp Lys His Pro Leu Ser Cys Ala 165 170 175

Cys Gly Arg Leu Cys Gly Glu Glu Ala Ala Asp Thr Gly Asp Asp Ile 180 185 190

Leu Pro His Glu Thr Gly Leu Gln Pro Gly Met Val Pro Leu Lys Tyr 195 200 205

Leu Leu Glu Glu Gly Val Trp Gly Ala Gly Val Gly Cys Gly Val Phe 210 215 220

Pro Ala Ile Ser Thr Ala Tyr Asp His Cys Asn Asn Leu Ser Pro Ser 225 230 235 235

Glu Glu His

<210> 174

<211> 147

<212> PRT

<213> Homo sapien

<400> 174

Met Ser Ser Glu Pro Pro Pro Pro Pro Gln Pro Pro Thr His Gln Ala 1 5 10 15

Ser Val Gly Leu Leu Asp Thr Pro Arg Ser Arg Glu Arg Ser Pro Ser 20 25 30

Pro Leu Arg Gly Asn Val Val Pro Ser Pro Leu Pro Thr Arg Arg Thr 35 40 45

Arg Thr Phe Ser Ala Thr Val Arg Ala Ser Gln Gly Pro Val Tyr Lys 50 55 60

Gly Val Cys Lys Cys Phe Cys Arg Ser Lys Gly His Gly Phe Ile Thr 65 70 75 80

Pro Ala Asp Gly Gly Pro Asp Ile Phe Leu His Ile Ser Asp Val Glu 85 90 95

Gly Glu Tyr Val Pro Val Glu Gly Asp Glu Val Thr Tyr Lys Met Cys 100 105 110

Ser Ile Pro Pro Lys Asn Glu Lys Leu Gln Ala Val Glu Val Val Ile 115 120 125

Thr His Leu Ala Pro Gly Thr Lys His Glu Thr Trp Ser Gly His Val

Ile Ser Ser 145

<210> 175

<211> 202

<212> PRT

<213> Homo sapien

<400> 175

Ser Ala Asp Gly Ala Pro Asp Ile Ser Pro Ser Leu Ser Gly Arg Cys 20 25 30

Gly Leu His Arg Ala Pro Ser Thr Lys Glu Ser Ala Asn Ala Ser Ala 35 40 45

Gly Pro Arg Ala Met Ala Ser Leu Pro Gln Leu Met Ala Ala Pro Thr 50 55 60

Ser Ser Cys Thr Ser Leu Met Trp Lys Gly Ser Met Ser Gln Trp Lys 65 70 75 80

Ala Thr Arg Ser Pro Ile Lys Cys Ala Pro Ser His Pro Arg Met Arg 85 90 95

Ser Cys Arg Pro Trp Arg Ser Ser Ser Leu Thr Trp His Gln Ala Pro 100 105 110

Ser Met Arg Pro Gly Leu Asp Met Ser Ser Ala Pro Arg Arg Trp Trp 115 120 125

Lys His Pro Leu Ser Cys Ala Cys Gly Arg Leu Cys Gly Glu Glu Ala 130 135 140

Ala Asp Thr Gly Asp Asp Ile Leu Pro His Glu Thr Gly Leu Gln Pro 145 150 155 160

Gly Met Val Pro Leu Lys Tyr Leu Leu Glu Glu Gly Val Trp Gly Ala 165 170 175

Gly Val Gly Cys Gly Val Phe Pro Ala Ile Ser Thr Ala Tyr Asp His 180 185 190

Cys Asn Asn Leu Ser Pro Ser Glu Glu His 195 200

<210> 176

<211> 138

<212> PRT

<213> Homo sapien

<400> 176

Met Ala Ser Leu Pro Gln Leu Met Ala Ala Pro Thr Ser Ser Cys Thr 1 5 10 15

Ser Leu Met Trp Lys Gly Ser Met Ser Gln Trp Lys Ala Thr Arg Ser 20 25 30

Pro Ile Lys Cys Ala Pro Ser His Pro Arg Met Arg Ser Cys Arg Pro 35 40 45

Trp Arg Ser Ser Ser Leu Thr Trp His Gln Ala Pro Ser Met Arg Pro 50 55 60

Gly Leu Asp Met Ser Ser Ala Pro Arg Arg Trp Trp Lys His Pro Leu 65 70 75 80

Ser Cys Ala Cys Gly Arg Leu Cys Gly Glu Glu Ala Ala Asp Thr Gly 85 90 95

Asp Asp Ile Leu Pro His Glu Thr Gly Leu Gln Pro Gly Met Val Pro 100 105 110

Leu Lys Tyr Leu Leu Glu Glu Gly Val Trp Gly Gly Arg Cys Gly Val

Trp Gly Val Pro Gly His Gln His Ser Leu 130 135

<210> 177

<211> 185

<212> PRT

<213> Homo sapien

<400> 177

Met Ser Ser Glu Pro Pro Pro Pro Pro Gln Pro Pro Thr His Gln Ala 1 5 10 15

Ser Val Gly Leu Leu Asp Thr Pro Arg Ser Arg Glu Arg Ser Pro Ser 20 25 30

Pro Ser Thr Lys Glu Ser Ala Asn Ala Ser Ala Gly Pro Arg Ala Met 35 40 45

Ala Ser Leu Pro Gln Leu Met Ala Ala Pro Thr Ser Ser Cys Thr Ser 50 60

Leu Met Trp Lys Gly Ser Met Ser Gln Trp Lys Ala Thr Arg Ser Pro 65 70 75 80

Ile Lys Cys Ala Pro Ser His Pro Arg Met Arg Ser Cys Arg Pro Trp 85 90 95

240

Arg Ser Ser Ser Leu Thr Trp His Gln Ala Pro Ser Met Arg Pro Gly 105 110

Leu Asp Met Ser Ser Ala Pro Arg Arg Trp Trp Lys His Pro Leu Ser 120 125

Cys Ala Cys Gly Arg Leu Cys Gly Glu Glu Ala Ala Asp Thr Gly Asp 130 135

Asp Ile Leu Pro His Glu Thr Gly Leu Gln Pro Gly Met Val Pro Leu 145 150 155

Lys Tyr Leu Leu Glu Glu Gly Val Trp Gly Gly Arg Cys Gly Val Trp 170

Gly Val Pro Gly His Gln His Ser Leu 180

<210> 178 <211> 265 <212> PRT <213> Homo sapien

<400> 178

Ser Phe Pro Pro Ala His Leu Phe Ser Ala Cys Arg Gly Ser Ser Ser

Arg Pro Pro Arg Cys Phe Cys Leu Trp Ala Gly Ala Leu Asp Gly Gly 25

Leu Ala Gly Arg Trp Gly Glu Ala Arg Gly Ala Ser His Ala Gly Ser 40

Arg Ala Thr Pro Arg Arg Ala Trp Pro Arg Gln Leu Trp Leu Glu Val 55

Gly Thr Ser Ala Met Ser Ser Glu Pro Pro Pro Pro Pro Gln Pro Pro

Thr His Gln Ala Ser Val Gly Leu Leu Asp Thr Pro Arg Ser Arg Glu 90

Arg Ser Pro Ser Pro Ser Thr Lys Glu Ser Ala Asn Ala Ser Ala Gly 105 100

Pro Arg Ala Met Ala Ser Leu Pro Gln Leu Met Ala Ala Pro Thr Ser 120 115

Ser Cys Thr Ser Leu Met Trp Lys Gly Ser Met Ser Gln Trp Lys Ala 130 135 140

Thr Arg Ser Pro Ile Lys Cys Ala Pro Ser His Pro Arg Met Arg Ser 145 150 155 160

Cys Arg Pro Trp Arg Ser Ser Leu Thr Trp His Gln Ala Pro Ser 165 170 175

Met Arg Pro Gly Leu Asp Met Ser Ser Ala Pro Arg Arg Trp Trp Lys 180 185 190

His Pro Leu Ser Cys Ala Cys Gly Arg Leu Cys Gly Glu Glu Ala Ala 195 200 205

Asp Thr Gly Asp Asp Ile Leu Pro His Glu Thr Gly Leu Gln Pro Gly 210 215 220

Met Val Pro Leu Lys Tyr Leu Leu Glu Glu Gly Val Trp Gly Ala Gly 225 230 235 240

Val Gly Cys Gly Val Phe Pro Ala Ile Ser Thr Ala Tyr Asp His Cys 245 250 255

Asn Asn Leu Ser Pro Ser Glu Glu His 260 265

<210> 179

<211> 201

<212> PRT

<213> Homo sapien

<400> 179

Met Cys Ser Thr Val Pro Trp Arg Arg Thr Leu Arg Glu Gly Gln Phe 1 5 10 15

Leu Pro Leu Pro Cys Gly Val Trp Leu Pro Ala Ala Ser Gly Arg
20 25 30

Val Arg Gly Val Ala Glu Phe Gly Ser Arg Trp Leu Ala Leu Lys Ser

Pro Trp Leu Trp Val Phe Phe Phe Glu Thr Glu Ser Cys Ser Val Ala 50 55 60

Gln Ala Gly Val Gln Trp Cys Asp Leu Ser Ser Leu Glu Pro Pro Pro 65 70 75 80

Pro Arg Phe Lys Gln Phe Ser Cys Leu Ser Leu Gln Val Asp Gly Ile 85 90 95

Thr Gly Ala Cys His His Ala Gln Leu Ile Phe Val Phe Val Leu Glu 100 105 110

Thr Gly Phe Pro His Val Gly Gln Ala Ser Leu Glu Leu Leu Thr Leu 115 120 125

Ser Asp Pro Pro Ala Ser Ala Ser Gln Ser Ala Gly Ile Ala Gly Val 130 135 140

Ser His Cys Ala Arg Pro Leu Ala Leu Gly Phe Leu Leu Thr Phe Leu 145 150 150

Leu Pro Ser His Lys Tyr Tyr Val Ser Gln Met Cys Arg Asp Pro Cys 165 170 175

Leu Val Leu Gly Thr Gln Arg Gly Pro Met Pro Ser Trp Ser Trp Gly 180 185 190

Arg Met Trp His Phe His Glu Glu Leu 195 200

<210> 180

<211> 159

<212> PRT

<213> Homo sapien

<400> 180

Met Ala Gln Gly Leu Ala Val Arg Glu Met Thr Gly Met Thr Lys Phe 1 5 10 15

Lys Pro Tyr Phe Ile Ser Ala Thr Ser Glu Ile Leu Ser Gln Lys Cys 20 25 30

Ile Asn Thr Asn Val Leu Phe Leu Ser Leu Ser Asp Asn His Gly His  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Ile Asp Pro Ser Leu Arg Leu Ile Trp Asp Leu Ala Phe Leu Gly Ser 50 55

Ser Tyr Val Met Trp Glu Met Thr Thr Gln Val Ser His Tyr Tyr Leu 65 70 75 80 Ala Gln Leu Thr Ser Val Arg Gln Trp Lys Thr Asn Asp Asp Thr Ile 85 90 95

Asp Phe Asp Tyr Thr Val Leu Leu His Glu Leu Ser Thr Gln Glu Ile 100 105 110

Ile Pro Cys Arg Ile His Leu Val Trp Tyr Pro Gly Lys Pro Leu Lys
115 120 125

Val Lys Tyr His Cys Gln Glu Leu Gln Thr Pro Glu Glu Ala Ser Gly 130 135 140

Thr Glu Glu Gly Ser Ala Val Val Pro Thr Glu Leu Ser Asn Phe 145 150 155

<210> 181

<211> 128

<212> PRT

<213> Homo sapien

<400> 181

Met Arg Ile His Asp Phe His Ile Leu Lys Gln Asn Thr Thr Lys Asn 1 5 10 15

Arg Glu Ala Glu Ile Glu Lys Ala Val Gly Asp Thr Arg His Pro Phe 20 25 30

Lys Thr Arg Met Tyr Cys Ile Val Ile Thr Leu Asn Thr Thr Ile Phe

Ile Thr Leu Thr Leu Phe Ser Pro Ser Arg Lys Thr Asn Asp Asp Thr 50 55 60

Ile Asp Phe Asp Tyr Thr Val Leu Leu His Glu Leu Ser Thr Gln Glu 65 70 75 80

Ile Ile Pro Cys Arg Ile His Leu Val Trp Tyr Pro Gly Lys Pro Leu 85 90 95

Lys Val Lys Tyr His Cys Gln Glu Leu Gln Thr Pro Glu Glu Ala Ser

Gly Thr Glu Glu Gly Ser Ala Val Val Pro Thr Glu Leu Ser Asn Phe 115 120 125

<210> 182 <211> 224 <212> PRT

<213> Homo sapien

<400> 182

Met Asp Gly Asp Gln Thr Glu Glu Arg Met Met Lys Met Met Val His 1 5 10 15

Gln Arg Pro Leu Pro Gln Pro Ala Leu Leu Pro Met Ser Ser Asn Thr 20 25 30

Phe Pro Ser Arg Ser Thr Lys Pro Ser Pro Met Asn Pro Leu Pro Ser 35 40 45

Ser His Met Pro Gly Ala Phe Ser Glu Ser Asn Ser Ser Phe Pro Gln 50 55 60

Ser Ala Ser Leu Pro Pro Tyr Phe Ser Gln Gly Pro Ser Asn Arg Pro 65 70 75 80

Pro Ile Arg Ala Glu Gly Arg Asn Phe Pro Leu Pro Leu Pro Asn Lys 85 90 95

Pro Arg Pro Pro Ser Pro Ala Glu Glu Glu Asn Ser Leu Asn Glu Glu 100 105 110

Trp Tyr Val Ser Tyr Ile Thr Arg Pro Glu Ala Glu Ala Ala Leu Arg 115 120 135

Lys Ile Asn Gln Asp Gly Thr Phe Leu Val Arg Asp Ser Ser Lys Lys 130 135 140

Thr Thr Thr Asn Pro Tyr Val Leu Met Val Leu Tyr Lys Asp Lys Val 145 150 155 160

Tyr Asn Ile Gln Ile Arg Tyr Gln Lys Glu Ser Gln Val Tyr Leu Leu 165 170 175

Gly Thr Gly Leu Arg Gly Lys Glu Asp Phe Leu Ser Val Ser Asp Ile 180 185 190

Ile Asp Tyr Phe Arg Lys Met Pro Leu Leu Leu Ile Asp Gly Lys Asn 195 200 205

Arg Gly Ser Arg Tyr Gln Cys Thr Leu Thr His Ala Ala Gly Tyr Pro 210 215 220

<210> 183 <211> 230 <212> PRT <213> Homo sapien

<400> 183

Met Phe Leu Ser Asp Ser Trp Gly Leu Pro Ile Ser Phe Ile Val Phe

Tyr Phe Cys Thr Val His Gln Arg Pro Leu Pro Gln Pro Ala Leu Leu 20

Pro Met Ser Ser Asn Thr Phe Pro Ser Arg Ser Thr Lys Pro Ser Pro

Met Asn Pro Leu Pro Ser Ser His Met Pro Gly Ala Phe Ser Glu Ser 55

Asn Ser Ser Phe Pro Gln Ser Ala Ser Leu Pro Pro Tyr Phe Ser Gln 70

Gly Pro Ser Asn Arg Pro Pro Ile Arg Ala Glu Gly Arg Asn Phe Pro 85

Leu Pro Leu Pro Asn Lys Pro Arg Pro Pro Ser Pro Ala Glu Glu 100 105

Asn Ser Leu Asn Glu Glu Trp Tyr Val Ser Tyr Ile Thr Arg Pro Glu 120 115

Ala Glu Ala Ala Leu Arg Lys Ile Asn Gln Asp Gly Thr Phe Leu Val 135 130

Arg Asp Ser Ser Lys Lys Thr Thr Thr Asn Pro Tyr Val Leu Met Val -155 145

Leu Tyr Lys Asp Lys Val Tyr Asn Ile Gln Ile Arg Tyr Gln Lys Glu 170 165

Ser Gln Val Tyr Leu Leu Gly Thr Gly Leu Arg Gly Lys Glu Asp Phe 180

Leu Ser Val Ser Asp Ile Ile Asp Tyr Phe Arg Lys Met Pro Leu Leu 205 200

246

Leu Ile Asp Gly Lys Asn Arg Gly Ser Arg Tyr Gln Cys Thr Leu Thr 210 220

His Ala Ala Gly Tyr Pro 225 230

<210> 184

<211> 225

<212> PRT

<213> Homo sapien

<400> 184

Gln Leu Gly Pro Ala Asn Leu Ile His Cys Phe Leu Leu Leu His Val 1 10 15

His Gln Arg Pro Leu Pro Gln Pro Ala Leu Leu Pro Met Ser Ser Asn 20 25 30

Thr Phe Pro Ser Arg Ser Thr Lys Pro Ser Pro Met Asn Pro Leu Pro 35 40 45

Ser Ser His Met Pro Gly Ala Phe Ser Glu Ser Asn Ser Ser Phe Pro 50 55 60

Gln Ser Ala Ser Leu Pro Pro Tyr Phe Ser Gln Gly Pro Ser Asn Arg 75 80

Pro Pro Ile Arg Ala Glu Gly Arg Asn Phe Pro Leu Pro Leu Pro Asn 85 90 95

Lys Pro Arg Pro Pro Ser Pro Ala Glu Glu Glu Asn Ser Leu Asn Glu 100 105 110

Glu Trp Tyr Val Ser Tyr Ile Thr Arg Pro Glu Ala Glu Ala Ala Leu 115 120 125

Arg Lys Ile Asn Gln Asp Gly Thr Phe Leu Val Arg Asp Ser Ser Lys 130 135 140

Lys Thr Thr Thr Asn Pro Tyr Val Leu Met Val Leu Tyr Lys Asp Lys 145 150 155 160

Val Tyr Asn Ile Gln Ile Arg Tyr Gln Lys Glu Ser Gln Val Tyr Leu 165 170 175

Leu Gly Thr Gly Leu Arg Gly Lys Glu Asp Phe Leu Ser Val Ser Asp 180 185 190

Ile Ile Asp Tyr Phe Arg Lys Met Pro Leu Leu Leu Ile Asp Gly Lys
195 200 205

Asn Arg Gly Ser Arg Tyr Gln Cys Thr Leu Thr His Ala Ala Gly Tyr 210 215 220

Pro 225

<210> 185

<211> 1085

<212> PRT

<213> Homo sapien

<400> 185

Met Ala Ala Ser Thr Gly Tyr Val Arg Leu Trp Gly Ala Ala Arg Cys 1 5 10 15

Trp Val Leu Arg Arg Pro Met Leu Ala Ala Gly Gly Arg Val Pro 20 25 30

Thr Ala Ala Gly Ala Trp Leu Leu Arg Gly Gln Arg Thr Cys Asp Ala 35 40 45

Ser Pro Pro Trp Ala Leu Trp Gly Arg Gly Pro Ala Ile Gly Gln 50 55 60

Trp Arg Gly Phe Trp Glu Ala Ser Ser Arg Gly Gly Gly Ala Phe Ser 65 70 75 80

Gly Glu Asp Ala Ser Glu Gly Gly Ala Glu Glu Gly Ala Gly Gly 85 90 95

Ala Gly Gly Ser Ala Gly Ala Gly Glu Gly Pro Val Ile Thr Ala Leu 100 105 110

Thr Pro Met Thr Ile Pro Asp Val Phe Pro His Leu Pro Leu Ile Ala 115 120 125

Ile Thr Arg Asn Pro Val Phe Pro Arg Phe Ile Lys Ile Ile Glu Val

Lys Asn Lys Lys Leu Val Glu Leu Leu Arg Arg Lys Val Arg Leu Ala 145 150 155 160

248

Gln Pro Tyr Val Gly Val Phe Leu Lys Arg Asp Asp Ser Asn Glu Ser 165 170 175

Asp Val Val Glu Ser Leu Asp Glu Ile Tyr His Thr Gly Thr Phe Ala 180 185 190

Gin Ile His Glu Met Gln Asp Leu Gly Asp Lys Leu Arg Met Ile Val 195 200 205

Met Gly His Arg Arg Val His Ile Ser Arg Gln Leu Glu Val Glu Pro 210 215 220

Glu Glu Pro Glu Ala Glu Asn Lys His Lys Pro Arg Arg Lys Ser Lys 225 230 235 240

Arg Gly Lys Lys Glu Ala Glu Asp Glu Leu Ser Ala Arg His Pro Ala 245 250 255

Glu Leu Ala Met Glu Pro Thr Pro Glu Leu Pro Ala Glu Val Leu Met 260 265 270

Val Glu Val Glu Asn Val Val His Glu Asp Phe Gln Val Thr Glu Glu 275 280 295

Val Lys Ala Leu Thr Ala Glu Ile Val Lys Thr Ile Arg Asp Ile Ile 290 295 300

Ala Leu Asn Pro Leu Tyr Arg Glu Ser Val Leu Gln Met Met Gln Ala 305 310 315 320

Gly Gln Arg Val Val Asp Asn Pro Ile Tyr Leu Ser Asp Met Gly Ala 325 330 335

Ala Leu Thr Gly Ala Glu Ser His Glu Leu Gln Asp Val Leu Glu Glu 340 345 350

Thr Asn Ile Pro Lys Arg Leu Tyr Lys Ala Leu Ser Leu Leu Lys Lys 355 360 365

Glu Phe Glu Leu Ser Lys Leu Gl<br/>n Gln Arg Leu Gly Arg Glu Val Glu 370 \$375\$ 380

Glu Lys Ile Lys Gln Thr His Arg Lys Tyr Leu Leu Gln Glu Gln Leu 385 390 395 400

Lys Ile Ile Lys Lys Glu Leu Gly Leu Glu Lys Asp Asp Lys Asp Ala

405 410 415

Ile Glu Glu Lys Phe Arg Glu Arg Leu Lys Glu Leu Val Val Pro Lys
420 425 430

His Val Met Asp Val Val Asp Glu Glu Leu Ser Lys Leu Gly Leu Leu 435 440 445

Asp Asn His Ser Ser Glu Phe Asn Val Thr Arg Asn Tyr Leu Asp Trp 450 455 460

Leu Thr Ser Ile Pro Trp Gly Lys Tyr Ser Asn Glu Asn Leu Asp Leu 465 470 475 480

Ala Arg Ala Gln Ala Val Leu Glu Glu Asp His Tyr Gly Met Glu Asp 485 490 495

Val Lys Lys Arg Ile Leu Glu Phe Ile Ala Val Ser Gln Leu Arg Gly 500 505 510

Ser Thr Gln Gly Lys Ile Leu Cys Phe Tyr Gly Pro Pro Gly Val Gly 515 520 525

Lys Thr Ser Ile Ala Arg Ser Ile Ala Arg Ala Leu Asn Arg Glu Tyr 530 540

Phe Arg Phe Ser Val Gly Gly Met Thr Asp Val Ala Glu Ile Lys Gly 545 550 555

His Arg Arg Thr Tyr Val Gly Ala Met Pro Gly Lys Ile Ile Gln Cys 565 570 575

Leu Lys Lys Thr Lys Thr Glu Asn Pro Leu Ile Leu Ile Asp Glu Val 580 585 590

Arg Ala Glu Arg Pro Gly Asp Pro Leu Thr Arg Gln Cys Arg Val Gln 595 600 605

Asp Glu Ala Glu Ala Phe Arg Val Leu Gly Ser Pro Ser Gln Leu Arg 610 615 620

Gly His Arg Arg. Asp Ala Gly Pro Asp Gly Thr Asp Gly Ser Leu Pro 625 630 635 640

Thr His Ser His Ala Gln Gln Ala Ala Arg Pro Arg Gln Gly Val Gly 645 650 655

- Ala Ala Leu Thr Arg Gly Leu Pro Ser Pro Asp Pro Gly Val Ser Pro 660 665 670
- Leu Gly Gly Leu Ala Arg Gly Thr Ala Arg Gly Thr Thr Cys Cys Leu 675 680 685
- Leu Pro Gln Val Asp Lys Ile Gly Arg Gly Tyr Gln Gly Asp Pro Ser 690 695 700
- Ser Ala Leu Leu Glu Leu Leu Asp Pro Glu Gln Asn Ala Asn Phe Leu 705 710 715 720
- Asp His Tyr Leu Asp Val Pro Val Asp Leu Ser Lys Val Gly Gly Leu 725 730 735
- Ser Gly Ala Trp Ala Cys Trp Gly Gly Tyr Ala Ala Ser Pro Pro Ala 740 745 750
- Pro Cys Arg Arg Pro Gln Val Leu Phe Ile Cys Thr Ala Asn Val Thr 755 760 765
- Asp Thr Ile Pro Glu Pro Leu Arg Asp Arg Met Glu Met Ile Asn Val 770 775 780
- Ser Gly Tyr Val Ala Gln Glu Lys Leu Ala Ile Ala Glu Arg Tyr Leu 785 790 795 800
- Val Pro Gln Ala Arg Ala Leu Cys Gly Leu Asp Glu Ser Lys Ala Lys 805 810 815
- Leu Ser Ser Asp Val Leu Thr Leu Leu Ile Lys Gln Tyr Cys Arg Glu 820 825 830
- Ser Gly Val Arg Asn Leu Gln Lys Gln Val Glu Lys Val Leu Arg Lys 835 840 845
- Ser Ala Tyr Lys Ile Val Ser Gly Glu Ala Glu Ser Val Glu Val Thr 850 855 860
- Pro Glu Asn Leu Gln Asp Phe Val Gly Lys Pro Val Phe Thr Val Glu 865 870 875 880
- Arg Met Tyr Asp Val Thr Pro Pro Gly Val Val Met Gly Leu Ala Trp 885 890 895

Thr Ala Met Gly Gly Ser Thr Leu Phe Val Glu Thr Ser Leu Arg Arg 900 905 910

Pro Gln Asp Lys Asp Ala Lys Gly Asp Lys Asp Gly Ser Leu Glu Val 915 920 925

Thr Gly Gln Leu Gly Glu Val Met Lys Glu Ser Ala Arg Ile Ala Tyr 930 935 940

Thr Phe Ala Arg Ala Phe Leu Met Gln His Ala Pro Ala Asn Asp Tyr 945 950 955 960

Leu Val Thr Ser His Ile His Leu His Val Pro Glu Gly Ala Thr Pro 965 970 975

Lys Asp Gly Pro Ser Ala Gly Cys Thr Ile Val Thr Ala Leu Leu Ser 980 985 990

Leu Ala Met Gly Arg Pro Val Arg Gln Asn Leu Ala Met Thr Gly Glu 995 1000 1005

Val Ser Leu Thr Gly Lys Ile Leu Pro Val Gly Gly Ile Lys Glu 1010 1015 1020

Lys Thr Ile Ala Ala Lys Arg Ala Gly Val Thr Cys Ile Val Leu 1025 1030 1035

Pro Ala Glu Asn Lys Lys Asp Phe Tyr Asp Leu Ala Ala Phe Ile 1040 1045 1050

Thr Glu Gly Leu Glu Val His Phe Val Glu His Tyr Arg Glu Ile 1055 1060 1065

Phe Asp Ile Ala Phe Pro Asp Glu Gln Ala Glu Ala Leu Ala Val 1070 1075 1080

Glu Arg 1085

<210> 186

<211> 668

<212> PRT

<213> Homo sapien

<400> 186

Asn Ser Ala Pro Ser Ser Pro Arg Arg Pro Ser Ser Leu Lys Arg Leu

. . .

252

1 5 10 15

Phe Arg Ala Thr Arg Pro Ser Gly Thr Glu Ala Arg Ala Gly Arg His 20 25 30

Val Arg Phe Ala Ala Ser Gly Asn Asp Ala Ser Cys Val Ser Arg Gln 35 40 45

Tyr Gly Arg Ala Met Ala Ala Ser Thr Gly Tyr Val Arg Leu Trp Gly
50 60

Ala Ala Arg Cys Trp Val Leu Arg Arg Pro Met Leu Ala Ala Ala Gly 65 70 75 80

Gly Arg Val Pro Thr Ala Ala Gly Ala Trp Leu Leu Arg Gly Gln Arg 85 90 95

Thr Cys Asp Ala Ser Pro Pro Trp Ala Leu Trp Gly Arg Gly Pro Ala 100 105 110

Ile Gly Gly Gln Trp Arg Gly Phe Trp Glu Ala Ser Ser Arg Gly Gly 115 120 125

Gly Ala Phe Ser Gly Gly Glu Asp Ala Ser Glu Gly Gly Ala Glu Glu 130 135 140

Gly Ala Gly Gly Ala Gly Gly Ser Ala Gly Ala Gly Glu Gly Pro Val 145 150 155 160

Ile Thr Ala Leu Thr Pro Met Thr Ile Pro Asp Val Phe Pro His Leu 165 170 175

Pro Leu Ile Ala Ile Thr Arg Asn Pro Val Phe Pro Arg Phe Ile Lys 180 185 190

Ile Ile Glu Val Lys Asn Lys Lys Leu Val Glu Leu Leu Arg Arg Lys
195 200 205

Val Arg Leu Ala Gln Pro Tyr Val Gly Val Phe Leu Lys Arg Asp Asp 210 215 220

Ser Asn Glu Ser Asp Val Val Glu Ser Leu Asp Glu Ile Tyr His Thr 225 230 235 240

Gly Thr Phe Ala Gln Ile His Glu Met Gln Asp Leu Gly Asp Lys Leu 245 250 255

Arg Met Ile Val Met Gly His Arg Arg Val His Ile Ser Arg Gln Leu 260 265 270

Glu Val Glu Pro Glu Glu Pro Glu Ala Glu Asn Lys His Lys Pro Arg 275 280 285

Arg Lys Ser Lys Arg Gly Lys Lys Glu Ala Glu Asp Glu Leu Ser Ala 290 295 300

Arg His Pro Ala Glu Leu Ala Met Glu Pro Thr Pro Glu Leu Pro Ala 305 310 315 320

Glu Val Leu Met Val Glu Val Glu As<br/>n Val Val His Glu Asp Phe Gl<br/>n 325 330 335

Val Thr Glu Glu Val Lys Ala Leu Thr Ala Glu Ile Val Lys Thr Ile 340 345 350

Arg Asp Ile Ile Ala Leu Asn Pro Leu Tyr Arg Glu Ser Val Leu Gln 355 360 365

Met Met Gln Ala Gly Gln Arg Val Val Asp Asn Pro Ile Tyr Leu Ser 370 380

Asp Met Gly Ala Ala Leu Thr Gly Ala Glu Ser His Glu Leu Gln Asp 385 390 395 400

Val Leu Glu Glu Thr Asn Ile Pro Lys Arg Leu Tyr Lys Ala Leu Ser 405 410 415

Leu Leu Lys Lys Glu Phe Glu Leu Ser Lys Leu Gln Gln Arg Leu Gly 420 425 430

Arg Glu Val Glu Glu Lys Ile Lys Gln Thr His Arg Lys Tyr Leu Leu 435 440 445

Gln Glu Gln Leu Lys Ile Ile Lys Lys Glu Leu Gly Leu Glu Lys Asp 450 455 460

Asp Lys Asp Ala Ile Glu Glu Lys Phe Arg Glu Arg Leu Lys Glu Leu 465 470 475 480

Val Val Pro Lys His Val Met Asp Val Val Asp Glu Glu Leu Ser Lys 485 490 495

Leu Gly Leu Leu Asp Asn His Ser Ser Glu Phe Asn Val Thr Arg Asn 500 505 510

Tyr Leu Asp Trp Leu Thr Ser Ile Pro Trp Gly Lys Tyr Ser Asn Glu 515 520 525

Asn Leu Asp Leu Ala Arg Ala Gln Ala Val Leu Glu Glu Asp His Tyr 530 540

Gly Met Glu Asp Val Lys Lys Arg Ile Leu Glu Phe Ile Ala Val Ser 545 550 555 560

Gln Leu Arg Gly Ser Thr Gln Gly Lys Ile Leu Cys Phe Tyr Gly Pro 565 570 575

Pro Gly Val Gly Lys Thr Ser Ile Ala Arg Ser Ile Ala Arg Ala Leu 580 585 590

Asn Arg Glu Tyr Phe Arg Phe Ser Val Gly Gly Met Thr Asp Val Ala 595 600 605

Glu Ile Lys Gly His Arg Arg Thr Tyr Val Gly Ala Met Pro Gly Lys 610 615 620

Ile Ile Gln Cys Leu Lys Lys Thr Lys Thr Glu Asn Pro Leu Ile Leu 625 630 635

Ile Asp Glu Val Arg Ala Glu Arg Pro Gly Asp Pro Leu Thr Arg Gln 645 650 655

Cys Arg Val Gln Asp Glu Ala Glu Pro Ser Gly Ser 660 665

<210> 187

<211> 771

<212> PRT

<213> Homo sapien

<400> 187

Met Glu Glu Arg Lys His Ser Leu Phe Phe Ser Pro Glu Ile Ala Tyr 1 5 10 15

Pro Lys Asp Lys Ala Leu Arg Gly Lys Glu Thr Gly Phe Gly Phe Ser 20 25 30

Phe Asp Ser Leu Gly Pro Val Ser Leu Gly Leu Val Ser Pro Pro Gly

255

35 40 45

Thr Ala Leu Leu His Leu Pro Gly Arg Pro Trp Pro Gly Pro Val Gly 50 55 60

Gly Gly Glu Gly Val Asp Gly Ala Gly Gly Trp Gly Ser Gln Gly His 65 70 75 80

Ala Pro Thr Gln Pro Thr Pro Cys Trp Ser Thr Ala Gly Pro Gln Arg 85 90 95

Pro Cys Pro Ile Leu Gly Ala Leu Thr Ala Glu Ile Val Lys Thr Ile 100 105 110

Arg Asp Ile Ile Ala Leu Asn Pro Leu Tyr Arg Glu Ser Val Leu Gln 115 120 125

Met Met Gln Ala Gly Gln Arg Val Val Asp Asn Pro Ile Tyr Leu Ser 130 135 140

Asp Met Gly Ala Ala Leu Thr Gly Ala Glu Ser His Glu Leu Gln Asp 145 150 155 160

Val Leu Glu Glu Thr Asn Ile Pro Lys Arg Leu Tyr Lys Ala Leu Ser 165 170 175

Leu Leu Lys Lys Glu Phe Glu Leu Ser Lys Leu Gln Gln Arg Leu Gly 180 185 190

Arg Glu Val Glu Glu Lys Ile Lys Gln Thr His Arg Lys Tyr Leu Leu 195 200 205

Gln Glu Gln Leu Lys Ile Ile Lys Lys Glu Leu Gly Leu Glu Lys Asp 210 215 220

Asp Lys Asp Ala Ile Glu Glu Lys Phe Arg Glu Arg Leu Lys Glu Leu 225 230 235 240

Val Val Pro Lys His Val Met Asp Val Val Asp Glu Glu Leu Ser Lys 245 250 255

Leu Gly Leu Leu Asp Asn His Ser Ser Glu Phe Asn Val Thr Arg Asn 260 265 270

Tyr Leu Asp Trp Leu Thr Ser Ile Pro Trp Gly Lys Tyr Ser Asn Glu 275 280 285

- Asn Leu Asp Leu Ala Arg Ala Gln Ala Val Leu Glu Glu Asp His Tyr 290 295 300
- Gly Met Glu Asp Val Lys Lys Arg Ile Leu Glu Phe Ile Ala Val Ser 305 310 315
- Gln Leu Arg Gly Ser Thr Gln Gly Lys Ile Leu Cys Phe Tyr Gly Pro 325 330 335
- Pro Gly Val Gly Lys Thr Ser Ile Ala Arg Ser Ile Ala Arg Ala Leu 340 345 350
- Asn Arg Glu Tyr Phe Arg Phe Ser Val Gly Gly Met Thr Asp Val Ala 355 360 365
- Glu Ile Lys Gly His Arg Arg Thr Tyr Val Gly Ala Met Pro Gly Lys 370 380
- Ile Ile Gln Cys Leu Lys Lys Thr Lys Thr Glu Asn Pro Leu Ile Leu 385 390 395 400
- Ile Asp Glu Val Asp Lys Ile Gly Arg Gly Tyr Gln Gly Asp Pro Ser 405 410 415
- Ser Ala Leu Leu Glu Leu Leu Asp Pro Glu Gln Asn Ala Asn Phe Leu 420 425 430
- Asp His Tyr Leu Asp Val Pro Val Asp Leu Ser Lys Val Leu Phe Ile 435 440 445
- Cys Thr Ala Asn Val Thr Asp Thr Ile Pro Glu Pro Leu Arg Asp Arg 450 455 460
- Met Glu Met Ile Asn Val Ser Gly Tyr Val Ala Gln Glu Lys Leu Ala 465 470 475 480
- Ile Ala Glu Arg Tyr Leu Val Pro Gln Ala Arg Ala Leu Cys Gly Leu 485 490 495
- Asp Glu Ser Lys Ala Lys Leu Ser Ser Asp Val Leu Thr Leu Leu Ile 500 505 510
- Lys Gln Tyr Cys Arg Glu Ser Gly Val Arg Asn Leu Gln Lys Gln Val 515 520 525

| Glu        | Lys<br>530 | Val        | Leu        | Arg        | Lys        | Ser<br>535 | Ala        | Tyr        | Lys        | Ile        | Val<br>540 | Ser        | Gly        | Glu        | Ala        |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu<br>545 | Ser        | Val        | Glu        | Val        | Thr<br>550 | Pro        | Glu        | Asn        | Leu        | Gln<br>555 | Asp        | Phe        | Val        | Gly        | Lys<br>560 |
| Pro        | Val        | Phe        | Thr        | Val<br>565 | Glu        | Arg        | Met        | Tyr        | Asp<br>570 | Val        | Thr        | Pro        | Pro        | Gly<br>575 | Val        |
| Val        | Met        | Gly        | Leu<br>580 | Ala        | Trp        | Thr        | Ala        | Met<br>585 | Gly        | Gly        | Ser        | Thr        | Leu<br>590 | Phe        | Val        |
| Glu        | Thr        | Ser<br>595 | Leu        | Arg        | Arg        | Pro        | Gln<br>600 | Asp        | Lys        | Asp        | Ala        | Lys<br>605 | Gly        | Asp        | Lys        |
| Asp        | Gly<br>610 | Ser        | Leu        | Glu        | Val        | Thr<br>615 | Gly        | Gln        | Leu        | Gly        | Glu<br>620 | Val        | Met        | Lys        | Glu        |
| Ser<br>625 | Ala        | Arg        | Ile        | Ala        | Tyr<br>630 | Thr        | Phe        | Ala        | Arg        | Ala<br>635 | Phe        | Leu        | Met        | Gln        | His<br>640 |
| Ala        | Pro        | Ala        | Asn        | Asp<br>645 | Tyr        | Leu        | Val        | Thr        | Ser<br>650 | His        | Ile        | His        | Leu        | His<br>655 | Val        |
| Pro        | Glu        | Gly        | Ala<br>660 | Thr        | Pro        | Lys        | qzA        | Gly<br>665 | Pro        | Ser        | Ala        | Gly        | Cys<br>670 | Thr        | Ile        |
| Val        | Thr        | Ala<br>675 | Leu        | Leu        | Ser        | Leu        | Ala<br>680 | Met        | Gly        | Arg        | Pro        | Val<br>685 | Arg        | Gln        | Asn        |
| Leu        | Ala<br>690 | Met        | Thr        | Gly        | Glu        | Val<br>695 | Ser        | Leu        | Thr        | Gly        | Lys<br>700 | Ile        | Leu        | Pro        | Val        |
| Gly<br>705 | Gly        | Ile        | Lys        | Glu        | Lys<br>710 | Thr        | Ile        | Ala        | Ala        | Lys<br>715 | Arg        | Ala        | Gly        | Val        | Thr<br>720 |
| Cys        | Ile        | Val        | Leu        | Pro<br>725 | Ala        | Glu        | Asn        | Lys        | Lys<br>730 | Asp        | Phe        | Tyr        | qzA        | Leu<br>735 | Ala        |
| Ala        | Phe        | Ile        | Thr        | Glu        | Gly        | Leu        | Glu        | Val        | His        | Phe        | Val        | Glu        | His        | Tyr        | Arg        |

740 745 750

Glu Ile Phe Asp Ile Ala Phe Pro Asp Glu Gln Ala Glu Ala Leu Ala

760

Val Glu Arg 770

<210> 188 <211> 848 <212> PRT <213> Homo sapien

<400> 188

Met Ala Ala Ser Thr Gly Tyr Val Arg Leu Trp Gly Ala Ala Arg Cys

Trp Val Leu Arg Arg Pro Met Leu Ala Ala Ala Gly Gly Arg Val Pro 25

Thr Ala Ala Gly Ala Trp Leu Leu Arg Gly Gln Arg Thr Cys Asp Ala 40

Ser Pro Pro Trp Ala Leu Trp Gly Arg Gly Pro Ala Ile Gly Gly Gln

Trp Arg Gly Phe Trp Glu Ala Ser Ser Arg Gly Gly Ala Phe Ser

Gly Gly Glu Asp Ala Ser Glu Gly Gly Ala Glu Glu Gly Ala Gly Gly

Ala Gly Gly Ser Ala Gly Ala Gly Glu Gly Pro Val Ile Thr Ala Leu

Thr Pro Met Thr Ile Pro Asp Val Phe Pro His Leu Pro Leu Ile Ala 120

Ile Thr Arg Asn Pro Val Phe Pro Arg Phe Ile Lys Ile Ile Glu Val 135

Lys Asn Lys Lys Leu Val Glu Leu Leu Arg Arg Lys Val Arg Leu Ala 145 150 155

Gln Pro Tyr Val Gly Val Phe Leu Lys Arg Asp Asp Ser Asn Glu Ser 165 170

Asp Val Val Glu Ser Leu Asp Glu Ile Tyr His Thr Gly Thr Phe Ala 185

Gln Ile His Glu Met Gln Asp Leu Gly Asp Lys Leu Arg Met Ile Val 200

| Met        | Gly<br>210 | His        | Arg        | Arg        | Val        | His<br>215 | Ile        | Ser        | Arg        | Gln        | Leu<br>220 | Glu        | Val        | Glu        | Pro        |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu<br>225 | Glu        | Pro        | Glu        | Ala        | Glu<br>230 | Asn        | Lys        | His        | Lys        | Pro<br>235 | Arg        | Arg        | Lys        | Ser        | Lys<br>240 |
| Arg        | Gly        | Lys        | Lys        | Glu<br>245 | Ala        | Glu        | Asp        | Glu        | Leu<br>250 | Ser        | Ala        | Arg        | His        | Pro<br>255 | Ala        |
| Glu        | Leu        | Ala        | Met<br>260 | Glu        | Pro        | Thr        | Pro        | Glu<br>265 | Leu        | Pro        | Ala        | Glu        | Val<br>270 | Leu        | Met        |
| Val        | Glu        | Val<br>275 | Glu        | Asn        | Val        | Val        | His<br>280 | Glu        | Asp        | Phe        | Gln        | Val<br>285 | Thr        | Glu        | Glu        |
| Val        | Lys<br>290 | Ala        | Leu        | Thr        | Ala        | Glu<br>295 | Ile        | Val        | Lys        | Thr        | Ile<br>300 | Arg        | Asp        | Ile        | Ile        |
| Ala<br>305 | Leu        | Asn        | Pro        | Leu        | Tyr<br>310 | Arg        | Glu        | Ser        | Val        | Leu<br>315 | Gln        | Met        | Met        | Gln        | Ala<br>320 |
| Gly        | Gln        | Arg        | Val        | Val<br>325 |            | Asn        | Pro        | Ile        | Tyr<br>330 | Leu        | Ser        | Asp        | Met        | Gly<br>335 | Ala        |
| Ala        | Leu        | Thr        | Gly<br>340 |            | Glu        | Ser        | His        | Glu<br>345 | Leu        | Gln        | Asp        | Val        | Leu<br>350 | Glu        | Glu        |
| Thr        | Asn        | Ile<br>355 | Pro        | Lys        | Arg        | Leu        | Tyr<br>360 | Lys        | Ala        | Leu        | Ser        | Leu<br>365 | Leu        | Lys        | Lys        |
| Glu        | Phe<br>370 | Glu        | Leu        | Ser        | Lys        | Leu<br>375 | Gln        | Gln        | Arg        | Leu        | Gly<br>380 |            | Glu        | Val        | Glu        |
| Glu<br>385 | -          | Ile        | Lys        | Gln        | Thr<br>390 | His        | Arg        | Lys        | Tyr        | Leu<br>395 |            | Gln        | Glu        | Gln        | Leu<br>400 |
| Lys        | Ile        | Ile        | Lys        | Lys<br>405 |            | Leu        | Gly        | Leu        | Glu<br>410 |            | Asp        | Asp        | Lys        | Asp<br>415 |            |
| Ile        | Glu        | Glu        | Lys<br>420 |            | Arg        | Glu        | Arg        | Leu<br>425 |            | Gly        | Leu        | . Val      | Val<br>430 |            | Lys        |
| His        | Val        | Met        | _          | Val        | . Val      | Asp        | Glu<br>440 |            | Leu        | Ser        | Lys        | Leu<br>445 | Gly        | Leu        | Leu        |

- Asp Asn His Ser Ser Glu Phe Asn Val Thr Arg Asn Tyr Leu Asp Trp 450 455 460
- Leu Thr Ser Ile Pro Trp Gly Lys Tyr Ser Asn Glu Asn Leu Asp Leu 465 470 475 480
- Ala Arg Ala Gl<br/>n Ala Val Leu Glu Glu Asp His Tyr Gly Met Glu Asp 495  $\phantom{\bigg|}$
- Val Lys Lys Arg Ile Leu Glu Phe Ile Ala Val Ser Gln Leu Arg Gly 500 505 510
- Ser Thr Gln Gly Lys Ile Leu Cys Phe Tyr Gly Pro Pro Gly Val Gly 515 520 525
- Lys Thr Ser Ile Ala Arg Ser Ile Ala Arg Ala Leu Asn Arg Glu Tyr 530 540
- Phe Arg Phe Ser Val Gly Gly Met Thr Asp Val Ala Glu Ile Lys Gly 545 550 555 560
- His Arg Arg Thr Tyr Val Gly Ala Met Pro Gly Lys Ile Ile Gln Cys 565 570 575
- Leu Lys Lys Thr Lys Thr Glu Asn Pro Leu Ile Leu Ile Asp Glu Val 580 585 590
- Asp Lys Ile Gly Arg Gly Tyr Gln Gly Asp Pro Ser Ser Ala Leu Leu 595 600 605
- Glu Leu Leu Asp Pro Glu Gln Asn Ala Asn Phe Leu Asp His Tyr Leu 610 615 620
- Asp Val Pro Val Asp Leu Ser Lys Val Leu Phe Ile Cys Thr Ala Asn 625 630 635 640
- Val Thr Asp Thr Ile Pro Glu Pro Leu Arg Asp Arg Met Glu Met Ile 645 650 655
- Asn Val Ser Gly Tyr Val Ala Gln Glu Lys Leu Ala Ile Ala Glu Arg 660 665 670
- Tyr Leu Val Pro Gln Ala Arg Ala Leu Cys Gly Leu Asp Glu Ser Lys 675 680 685

261

Ala Lys Leu Ser Ser Asp Val Leu Thr Leu Leu Ile Lys Gln Tyr Cys 690 695 700

Arg Glu Ser Gly Val Arg Asn Leu Gln Lys Gln Val Glu Lys Val Leu 705 710 715 720

Arg Lys Ser Ala Tyr Lys Ile Val Ser Gly Glu Ala Glu Ser Val Glu 725 730 735

Val Thr Pro Glu Asn Leu Gln Asp Phe Val Gly Lys Pro Val Phe Thr 740 745 750

Val Glu Arg Met Tyr Asp Val Thr Pro Pro Gly Val Val Met Gly Leu 755 760 765

Ala Trp Thr Ala Met Gly Glu Arg Gly Gly Gly Arg Arg Pro Gln Ser 770 775 780

His Ser His Phe Tyr Pro Arg Thr Ser Arg Ser His Leu Val His Leu 785 790 795 800

Cys Ser Gly Pro Gln Val Ala Leu Asn Gly Leu Trp Arg Gly Val Val 805 810 815

Cys Trp Glu Pro Arg Gly Ser Gly Leu Arg Glu Gly Lys Ala Val Thr 820 825 830

His Gly Gly Gly Ser Thr Pro Trp Ala Leu Trp Ile Trp Val Pro Leu 835 840 845

<210> 189

<211> 124

<212> PRT

<213> Homo sapien

<400> 189

Met Val Phe Leu His Val Gly Gln Ala Gly Leu Glu Leu Pro Thr Ser 1 5 10 15

Gly Asp Pro Pro Thr Ser Ala Ser Gln Ser Ala Gly Met Thr Glu Leu 20 25 30

Glu Leu Gly Pro Ser Pro Arg Leu Gln Pro Ile Arg Arg His Pro Lys

Glu Leu Pro Gln Tyr Gly Gly Pro Gly Lys Asp Ile Phe Glu Asp Gln 50 55 60

Leu Tyr Leu Pro Val His Ser Asp Gly Ile Ser Val His Gln Met Phe 65 70 75 80

Thr Met Ala Thr Ala Glu His Arg Ser Asn Ser Ser Ile Ala Gly Lys 85 90 95

Met Leu Thr Lys Val Glu Lys Asn His Glu Lys Glu Lys Ser Gln His 100 105 110

Leu Glu Gly Ser Ala Ser Ser Ser Leu Ser Ser Asp 115 120

<210> 190

<211> 296

<212> PRT

<213> Homo sapien

<400> 190

Met Ser Thr Glu Arg Thr Ser Trp Thr Ser Leu Ser Thr Ile Gln Lys
1 5 10 15

Leu Tyr Arg Arg Tyr Arg Glu Ser Arg Glu Glu Arg Leu Thr Phe Val 35 40 45

Gly Glu Asp Asp Ile Glu Ile Glu Met Arg Val Pro Gln Glu Ala Val 50 55 60

Lys Leu Ile Ile Gly Arg Gln Gly Ala Asn Ile Lys Gln Leu Arg Lys 65 70 75 80

Gln Thr Gly Ala Arg Ile Asp Val Asp Thr Glu Asp Val Gly Asp Glu 85 90 95

Arg Val Leu Leu Ile Ser Gly Phe Pro Val Gln Val Cys Lys Ala Lys
100 105 110

Ala Ala Ile His Gln Ile Leu Thr Glu Asn Thr Pro Val Ser Glu Gln 115 120 125

Leu Ser Val Pro Gln Arg Ser Val Gly Arg Ile Ile Gly Arg Gly Gly 130 135 140

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Glu Thr Ile Arg Ser Ile Cys Lys Ala Ser Gly Ala Lys Ile Thr Cys 150 155

Asp Lys Glu Ser Glu Gly Thr Leu Leu Leu Ser Arg Leu Ile Lys Ile 170 165

Ser Gly Thr Gln Lys Glu Val Ala Ala Ala Lys His Leu Ile Leu Glu 185

Lys Val Ser Glu Asp Glu Glu Leu Arg Lys Arg Ile Ala His Ser Ala

Glu Thr Arg Val Pro Arg Lys Gln Pro Ile Ser Val Arg Arg Glu Asp 215

Met Thr Glu Pro Gly Gly Ala Gly Glu Pro Ala Leu Trp Lys Asn Thr 235 230

Ser Ser Ser Met Glu Pro Thr Ala Pro Leu Val Thr Pro Pro Pro Lys 250 245

Gly Gly Gly Asp Met Ala Val Val Ser Lys Glu Gly Ser Trp Glu 265 260

Lys Pro Ser Asp Asp Ser Phe Gln Lys Ser Glu Ala Gln Ala Ile Pro 280

Glu Met Pro Met Phe Glu Ser Met 290

<210> 191

<211> 195 <212> PRT

<213> Homo sapien

<400> 191

Ala Arg Tyr Glu Ala Trp Gly Glu Ser Ala Glu Ala His Val Leu Glu

Gly Pro Asp Thr. Asn Thr Thr Ile Ile Gln Leu Gln Pro Leu Gln Glu 25 20

Pro Glu Ser Trp Ala Arg Thr Gln Ser Gly Leu Gln Ser Tyr Leu Leu 40 35

Gln Phe His Gly Leu Val Arg Leu Val His Gln Glu Arg Thr Leu Ala 55 50

Phe Pro Leu Thr Ile Arg Cys Phe Leu Gly Cys Glu Leu Pro Pro Glu 65 70

Gly Ser Arg Ala His Val Phe Phe Glu Val Ala Val Asn Gly Ser Ser

Phe Val Ser Phe Arg Pro Glu Arg Ala Leu Trp Gln Ala Asp Thr Gln 105

Val Thr Ser Gly Val Val Thr Phe Thr Leu Gln Gln Leu Asn Ala Tyr 120

Asn Arg Thr Arg Tyr Glu Leu Arg Glu Phe Leu Glu Asp Thr Cys Val 135

Gln Tyr Val Gln Lys His Ile Ser Ala Glu Asn Thr Lys Gly Ser Gln 150

Thr Ser Arg Ser Tyr Thr Ser Leu Val Leu Gly Val Leu Val Gly Ser 170 175

Phe Ile Ile Ala Gly Val Ala Val Gly Ile Phe Leu Cys Thr Gly Gly 185

Arg Arg Cys 195

<210> 192

<211> 194 <212> PRT

<213> Homo sapien

<220>

<221> MISC\_FEATURE

<222> (6)..(7) <223> X=any amino acid

<400> 192

Leu Gly Thr Gly Arg Xaa Xaa Ser Ala Glu Ala His Val Leu Glu Gly 10 5

Pro Asp Thr Asn Thr Thr Ile Ile Gln Leu Gln Pro Leu Gln Glu Pro 20

Glu Ser Trp Ala Arg Thr Gln Ser Gly Leu Gln Ser Tyr Leu Leu Gln 35 40

Phe His Gly Leu Val Arg Leu Val His Gln Glu Arg Thr Leu Ala Phe 50 55 60

Pro Leu Thr Ile Arg Cys Phe Leu Gly Cys Glu Leu Pro Pro Glu Gly 65 70 75 80

Ser Arg Ala His Val Phe Phe Glu Val Ala Val Asn Gly Ser Ser Phe 85 90 95

Val Ser Phe Arg Pro Glu Arg Ala Leu Trp Gln Ala Asp Thr Gln Val 100 105 110

Thr Ser Gly Val Val Thr Phe Thr Leu Gln Gln Leu Asn Ala Tyr Asn 115 120 125

Arg Thr Arg Tyr Glu Leu Arg Glu Phe Leu Glu Asp Thr Cys Val Gln 130 135 140

Tyr Val Gln Lys His Ile Ser Ala Glu Asn Thr Lys Gly Ser Gln Thr 145 150 155 160

Ser Arg Ser Tyr Thr Ser Leu Val Leu Gly Val Leu Val Gly Ser Phe 165 170 175

Ile Ile Ala Gly Val Ala Val Gly Ile Phe Leu Cys Thr Gly Gly Arg 180 185 190

Arg Cys

<210> 193

<211> 132

<212> PRT

<213> Homo sapien

<400> 193

Met Asn Tyr Thr Gln Arg Glu Leu Gln Met Ala Ala Pro Thr His Leu 1 5 10 15

Asn His Thr Val Val Gly Thr Pro Cys Gly Asn Gln Thr Leu Ala Thr
20 25 30

Thr Arg Arg Lys His Leu Ala Trp Arg Glu Arg Arg Pro Ala His Thr 35 40 45

Thr Pro Ala Arg Ala Arg Asp Gly Asn Pro Asn Ile Gly Val Gly Ala 50 55 60

Ala Asp Lys Pro Pro Ser Leu Leu Asn His Ala Arg Arg Ser Ser Leu 65 70 75 80

Pro Asn Arg Pro Pro Arg Ser Thr Gly Gly Asp Glu Ser Leu Ile Thr 85 90 95

His Asn Pro Ser Tyr Ser His Gly Arg Arg Ala Ile Leu His Ala Cys 100 105 110

Val Val Pro His His Thr Glu Arg Arg Val Ala Ser Ile Ile Cys Arg 115 120 125

Pro Gly Pro Arg 130

<210> 194

.<211> 199

<212> PRT

<213> Homo sapien

<400> 194

Ile Ile Thr Ile Lys Leu Phe Lys Lys Lys Lys Lys Lys Gln Thr Lys

1 10 15

Asn Ile Lys Thr Lys Lys Gln Lys Glu Lys Lys Lys Lys Gln Asn Arg 20 25 30

Gly Ala Gly Ala Pro Gln Lys Lys Ser Arg Pro Arg Gly Ala Ala Pro 35 40 45

Ile Thr Ala Arg Pro Pro Gly Phe Leu Val Ser Thr Gln Gln Gly Gly 50 55 60

Ala Pro His Lys Asn Glu Arg Ala His Arg Ser His Asn Tyr Thr Gln 65 70 75 80

Arg Glu Leu Arg Trp Pro Arg Pro His Thr Ser Thr Thr Gln Ser Trp 85 90 95

Val His Pro Ala Ala Thr Lys Arg Trp Pro Gln Pro Gly Gly Asn Ile 100 105 110

Leu Arg Gly Glu Lys Gly Asp Pro Leu Thr Leu Pro Gln Arg Glu Arg 115 120 125 Gly Thr Ala Thr Gln Thr Ser Val Trp Gly Pro Pro Thr Asn Leu Pro 135 140

His Ser Ser Thr Met Arg Asp Asp Leu His Tyr Gln Ile Ala Pro Leu 145 . 150 155

Ala Arg Arg Ala Ala Thr Asn Arg Ser Ser His Thr Ile His Leu Ile 170

His Thr Ala Val Ala Leu Ser Ser Thr Arg Val Trp Cys Leu Ile Thr 180 185

Pro Asn Asp Gly Ser His Gln 195

<210> 195 <211> 259 <212> PRT <213> Homo sapien

<400> 195

Met Glu Gly Arg Arg Gln Ala Arg Arg Leu Arg Gln Leu Ala Gly Ala

Gly Ala Gln Gly Gly Ser Pro Ile Val Glu Ala Ala Glu Asn Cys His

Gly Ala Ala Ser Val Gln Arg Ala Ser Leu Ile Tyr Ser Gly Arg Val 40

Cys Ile Trp Gly Gly Val Gln Gly Ala Asp Lys Asp Arg Arg Ala Pro

Gly Ala Ala Ala Gly Gly Ala Lys Thr Gly Thr Arg Met Ser Ala Pro

Gln Arg Pro Trp Ala Leu Ala Ala Gly Ala Arg Arg Thr Pro Arg Glu

Ala Gly Ile Ser Lys Ala Val Arg Pro Glu Ala Arg Pro Arg Leu Arg .. 100 ... 105

Thr Lys Thr Glu Leu Glu Asp Leu Gln Lys Lys Pro Pro Pro Tyr Leu 115 120

268

Arg Asn Leu Ser Ser Asp Asp Ala Asn Val Leu Val Trp His Ala Leu 130 135 140

Leu Leu Pro Asp Gln Pro Pro Tyr His Leu Lys Ala Phe Asn Leu Arg 145 150 155 160

Ile Ser Phe Pro Pro Glu Tyr Pro Phe Lys Pro Pro Met Ile Lys Phe 165 170 175

Thr Thr Lys Ile Tyr His Pro Asn Val Asp Glu Asn Gly Gln Ile Cys 180 185 190

Leu Pro Ile Ile Ser Ser Glu Asn Trp Lys Pro Cys Thr Lys Thr Cys 195 200 205

Gln Val Leu Glu Ala Leu Asn Val Leu Val Asn Arg Pro Asn Ile Arg 210 215 220

Glu Pro Leu Arg Met Asp Leu Ala Asp Leu Leu Thr Gln Asn Pro Glu 225 230 235 240

Leu Phe Arg Lys Asn Ala Glu Glu Phe Thr Leu Arg Phe Gly Val Asp 245 250 255

Arg Pro Ser

<210> 196

<211> 156

<212> PRT

<213> Homo sapien

<400> 196

Met Arg Arg Trp Thr Trp Lys Ala Ser Trp Ser Ser Glu Leu Glu Asp 1 5 10 15

Leu Gln Lys Lys Pro Pro Pro Tyr Leu Arg Asn Leu Ser Ser Asp Asp 20 25 30

Ala Asn Val Leu Val Trp His Ala Leu Leu Leu Pro Asp Gln Pro Pro 35 40 45

Tyr His Leu Lys Ala Phe Asn Leu Arg Ile Ser Phe Pro Pro Glu Tyr 50 55 60

Pro Phe Lys Pro Pro Met Ile Lys Phe Thr Thr Lys Ile Tyr His Pro 65 70 75 80

Asn Val Asp Glu Asn Gly Gln Ile Cys Leu Pro Ile Ile Ser Ser Glu 85 90 95

Asn Trp Lys Pro Cys Thr Lys Thr Cys Gln Val Leu Glu Ala Leu Asn 100 105 110

Val Leu Val Asn Arg Pro Asn Ile Arg Glu Pro Leu Arg Met Asp Leu 115 120 125

Ala Asp Leu Leu Thr Gln Asn Pro Glu Leu Phe Arg Lys Asn Ala Glu 130 135 140

Glu Phe Thr Leu Arg Phe Gly Val Asp Arg Pro Ser 145 150 155

<210> 197

<211> 168

<212> PRT

<213> Homo sapien

<400> 197

Asn Leu Gly Gln Ser Ser Leu Ser Ile Leu Trp Arg Lys Arg Cys Gly
1 5 10 15

Gly Gly Pro Gly Arg Pro Pro Gly Glu Leu Glu Asp Leu Gln Lys Lys 20 25 30

Pro Pro Pro Tyr Leu Arg Asn Leu Ser Ser Asp Asp Ala Asn Val Leu 35 40

Val Trp His Ala Leu Leu Pro Asp Gln Pro Pro Tyr His Leu Lys 50 55 60

Ala Phe Asn Leu Arg Ile Ser Phe Pro Pro Glu Tyr Pro Phe Lys Pro 65 70 75 80

Pro Met Ile Lys Phe Thr Thr Lys Ile Tyr His Pro Asn Val Asp Glu 85 90 95

Asn Gly Gln Ile Cys Leu Pro Ile Ile Ser Ser Glu Asn Trp Lys Pro 100 105 110

Cys Thr Lys Thr Cys Gln Val Leu Glu Ala Leu Asn Val Leu Val Asn 115 120 125

Arg Pro Asn Ile Arg Glu Pro Leu Arg Met Asp Leu Ala Aso Leu Leu 130 135

Thr Gln Asn Pro Glu Leu Phe Arg Lys Asn Ala Glu Glu Phe Thr Leu 155

Arg Phe Gly Val Asp Arg Pro Ser 165

<210> 198

<211> 137

<212> PRT

<213> Homo sapien

<400> 198

Met Glu Gly Arg Arg Gln Gly Asn Leu Ser Ser Asp Asp Ala Asn Val

Leu Val Trp His Ala Leu Leu Leu Pro Asp Gln Pro Pro Tyr His Leu

Lys Ala Phe Asn Leu Arg Ile Ser Phe Pro Pro Glu Tyr Pro Phe Lys

Pro Pro Met Ile Lys Phe Thr Thr Lys Ile Tyr His Pro Asn Val Asp

Glu Asn Gly Gln Ile Cys Leu Pro Ile Ile Ser Ser Glu Asn Trp Lys 65 70

Pro Cys Thr Lys Thr Cys Gln Val Leu Glu Ala Leu Asn Val Leu Val 90

Asn Arg Pro Asn Ile Arg Glu Pro Leu Arg Met Asp Leu Ala Asp Leu 100 105

Leu Thr Gln Asn Pro Glu Leu Phe Arg Lys Asn Ala Glu Glu Phe Thr

Leu Arg Phe Gly Val Asp Arg Pro Ser 135

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<210> 199

<211> 237

<212> PRT

<213> Homo sapien

<400> 199

Met Met Ala Ser Met Arg Val Val Lys Glu Leu Glu Asp Leu Gln Lys 10

Lys Pro Pro Pro Tyr Leu Arg Asn Leu Ser Ser Asp Asp Ala Asn Val

Leu Val Trp His Ala Leu Leu Leu Pro Glu Ala Glu Val Ala Val Ser 40

Arg Asp His Ala Ile Ala Leu Gln Pro Gly Gln Gln Ser Glu Thr Pro

Ser Gln Lys Lys Lys Lys Glu Ala Trp His Gln His Leu Leu Leu

Met Arg Pro Ser Gly Ser Phe His Ser Trp Trp Lys Ala Lys Gly Ser 90.

His Val Tyr Arg Ser His Ala Arg Glu Glu Val Lys Glu Arg Glu Ser 105

Glu Gln Val Pro Gly Ser Ser Lys Gln Pro Ala Phe Ser Asp Gln Pro

Pro Tyr His Leu Lys Ala Phe Asn Leu Arg Ile Ser Phe Pro Pro Glu

Tyr Pro Phe Lys Pro Pro Met Ile Lys Phe Thr Thr Lys Ile Tyr His 155 150

Pro Asn Val Asp Glu Asn Gly Gln Ile Cys Leu Pro Ile Ile Ser Ser 165

Glu Asn Trp Lys Pro Cys Thr Lys Thr Cys Gln Val Leu Glu Ala Leu 180

Asn Val Leu Val Asn Arg Pro Asn Ile Arg Glu Pro Leu Arg Met Asp 200

Leu Ala Asp Leu Leu Thr Gln Asn Pro Glu Leu Phe Arg Lys Asn Ala 215 220

Glu Glu Phe Thr Leu Arg Phe Gly Val Asp Arg Pro Ser 230

<210> 200

<211> 156

<212> PRT

<213> Homo sapien

<400> 200

Gly Pro Gln Glu Ala Ser Thr His Gly Gly Arg Gln Arg Gly Ala Thr 10

Cys Thr Asp His Ile Ala Arg Glu Glu Val Lys Glu Arg Glu Ser Glu 25

Gln Val Pro Gly Ser Ser Lys Gln Pro Ala Phe Ser Asp Gln Pro Pro 40

Tyr His Leu Lys Ala Phe Asn Leu Arg Ile Ser Phe Pro Pro Glu Tyr 55

Pro Phe Lys Pro Pro Met Ile Lys Phe Thr Thr Lys Ile Tyr His Pro 70

Asn Val Asp Glu Asn Gly Gln Ile Cys Leu Pro Ile Ile Ser Ser Glu

Asn Trp Lys Pro Cys Thr Lys Thr Cys Gln Val Leu Glu Ala Leu Asn 105 100

Val Leu Val Asn Arg Pro Asn Ile Arg Glu Pro Leu Arg Met Asp Leu 120

Ala Asp Leu Leu Thr Gln Asn Pro Glu Leu Phe Arg Lys Asn Ala Glu 135

Glu Phe Thr Leu Arg Phe Gly Val Asp Arg Pro Ser

<210> 201

<211> 88

<212> PRT

<213> Homo sapien

<400> 201

Met Val Gln Ala Gly Pro Ser Ser Cys Ser Ile Ser Gly Asp Pro Gly

Leu Pro Arg Arg Trp Arg Pro Ala Gln Val Val Arg Pro Gly Arg Leu 25 20

273

Arg Ile Arg Gly Trp Ser Arg Arg Ile Pro Lys Ala Glu Val Gly Ser

Pro Gly Asp Ser Gln Leu Leu Ser Leu Trp Arg Arg Gly Pro Val Thr 55

Glu Ala Pro Phe Ser Asn Pro Gly Ala Ala Phe Gln Arg Leu Asn Phe 70

Ser Asn His Cys Phe Asn Ser Phe 85

<210> 202

<211> 40

<212> PRT

<213> Homo sapien

<400> 202

Met Glu Lys Gly Val Gly Gln Pro Arg Gly Arg Arg Ile Tyr

Asn Ile Phe Phe Arg His Arg Cys Tyr Arg Lys Met Cys Glu Arg Ser 25

Gly Cys Ala Ala Arg Thr Gly Ala

<210> 203 <211> 60 <212> PRT <213> Homo sapien

<400> 203

Gly Pro Glu Lys Trp Arg Arg Gly Trp Gly Asp Ser His Val Ala Ala 10

Gly Gly Phe Thr Thr Phe Ser Phe Ala Ile Asp Val Ile Ala Lys Cys 25

Val Arg Glu Ala Ala Ala Gln Pro Gly Arg Glu Arg Glu Gly Ala Gly

Gln Arg Phe Pro Pro Thr Gly Asn Leu Met Gly Leu 50 55

<210> 204

<211> 213

<212> PRT

<213> Homo sapien

<220>

<221> MISC\_FEATURE

<222> (5)..(5)

<223> X=any amino acid

<400> 204

Met Pro Gln Asn Xaa Gly Val Ile Gly Leu Arg His His Phe Ala Ile

His Tyr Pro Ala Gly Gly Gly Leu Trp Asp Gly Leu His Gly Val Ala

Ala Val Gln Gly Ile Thr Lys Ile Lys Val Leu Ala Ile Tyr Ser Phe 40

Cys Ser Gln Ile Cys Asp Pro Arg Thr Thr Gly Ala Phe Trp Gln Thr 50

Trp Lys Asp Phe Glu Val Arg His Gly Asn Glu Asp Thr Ile Lys Glu

Met Leu Arg Ile Arg Arg Ser Val Gln Ala Thr Tyr Asn Thr Gln Val

Asn Phe Met Ala Ser Gln Met Leu Lys Val Ser Gly Ser Ala Thr Gly

Thr Val Ser Asp Leu Ala Pro Gly Gln Ser Gly Met Asp Asp Met Lys 120 115

Leu Leu Glu Gln Arg Ala Glu Gln Leu Ala Ala Glu Ala Glu Arg Asp 135 130

Gln Pro Leu Arg Ala Gln Ser Lys Ile Leu Phe Val Arg Ser Asp Ala 150

Ser Arg Glu Glu Leu Ala Glu Leu Ala Gln Gln Val Asn Pro Glu Glu 170 165

Ile Gln Leu Gly Glu Asp Glu Asp Glu Asp Glu Met Asp Leu Glu Pro 135 180

Asn Glu Val Arg Leu Glu Gln Gln Ser Val Pro Ala Ala Val Phe Gly 200 195

| ser                          | 210        | гÀг                     | GIU        | Asp        |            |            |            |            |            |            |            |            |            |            |            |
|------------------------------|------------|-------------------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| <210<br><211<br><212<br><213 | > 4<br>> E | 05<br>58<br>PRT<br>Iomo | sapi       | en         |            |            |            |            |            |            |            |            |            |            |            |
| <400                         | )> 2       | 05                      |            |            |            |            |            |            |            |            |            |            |            |            |            |
| Met<br>1                     | Ile        | Asp                     | His        | Tyr<br>5   | Arg        | Gly        | Met        | Gly        | Pro<br>10  | Leu        | Met        | Leu        | Leu        | Glu<br>15  | Arg        |
| Arg                          | Ser        | Val                     | Met<br>20  | Asp        | Arg        | Gly        | Arg        | Gly<br>25  | Arg        | Tyr        | Gln        | Tyr        | Ser<br>30  | Pro        | Gln        |
| Asn                          | Gln        | His<br>35               | Val        | Glu        | Gln        | Gln        | Pro<br>40  | His        | Tyr        | Thr        | His        | Lys<br>45  | Pro        | Thr        | Leu        |
| Glu                          | Tyr<br>50  | Ser                     | Pro        | Phe        | Pro        | Ile<br>55  | Pro        | Pro        | Gln        | Ser        | Pro<br>60  | Ala        | Tyr        | Glu        | Pro        |
| Asn<br>65                    | Leu        | Phe                     | Asp        | Gly        | Pro<br>70  | Glu        | Ser        | Gln        | Phe        | Cys<br>75  | Pro        | Asn        | Gln        | Ser        | Leu<br>80  |
| Val                          | Ser        | Leu                     | Leu        | Gly<br>85  | Asp        | Gln        | Arg        | Glu        | Ser<br>90  | Glu        | Asn        | Ile        | Ala        | Asn<br>95  | Pro        |
| Met                          | Gln        | Thr                     | Ser<br>100 | Ser        | Ser        | Val        | Gln        | Gln<br>105 | Gln        | Asn        | Asp        | Ala        | His<br>110 | Leu        | His        |
| Ser                          | Phe        | Ser<br>115              | Met        | Met        | Pro        | Ser        | Ser<br>120 | Ala        | Cys        | Glu        | Ala        | Met<br>125 | Val        | Gly        | His        |
| Glu                          | Met<br>130 | Ala                     | Ser        | Asp        | Ser        | Ser<br>135 | Asn        | Thr        | Ser        | Leu        | Pro<br>140 | Phe        | Ser        | Asn        | Met        |
| Gly<br>145                   | Asn        | Pro                     | Met        |            | Thr<br>150 | Thr        | Gln        | Leu        | Gly        | Lys<br>155 | Ser        | Leu        | Phe        | Gln        | Trp<br>160 |
| Gln                          | Val        |                         | Gln        | Glu<br>165 |            | Ser        |            | Leu        | Ala<br>170 | Asn<br>    | Ile        | Ser        | Gln        | Asp<br>175 | Gln        |
| Phe                          | Leu        | Ser                     | Lys<br>180 | Asp        | Ala        | Asp        | Gly        | Asp<br>185 | Thr        | Phe        | Leu        | His        | Ile<br>190 | Ala        | Val        |

276

Ala Gln Gly Arg Arg Ala Leu Ser Tyr Val Leu Ala Arg Lys Met Asn 195 200 205

Ala Leu His Met Leu Asp Ile Lys Glu His Asn Gly Gln Ser Ala Phe 210 220

Gln Val Ala Val Ala Ala Asn Gln His Leu Ile Val Gln Asp Leu Val 225 230 235 240

Asn Ile Gly Ala Gln Val Asn Thr Thr Asp Cys Trp Gly Arg Thr Pro 245 250 255

Leu His Val Cys Ala Glu Lys Gly His Ser Gln Val Leu Gln Ala Ile 260 265 270

Gln Lys Gly Ala Val Gly Ser Asn Gln Phe Val Asp Leu Glu Ala Thr 275 280 285

Asn Tyr Asp Gly Leu Thr Pro Leu His Cys Ala Val Ile Ala His Asn 290 295 300

Ala Val Val His Glu Leu Gln Arg Asn Gln Gln Pro His Ser Pro Glu 305 310 315 320

Val Gln Glu Leu Leu Lys Asn Lys Ser Leu Val Asp Thr Ile Lys 325 330 335

Cys Leu Ile Gln Met Gly Ala Ala Val Glu Ala Lys Asp Arg Lys Ser 340 345 350

Gly Arg Thr Ala Leu His Leu Ala Ala Glu Glu Ala Asn Leu Glu Leu 355 360 365

Ile Arg Leu Phe Leu Glu Leu Pro Ser Cys Leu Ser Phe Val Asn Ala 370 380

Lys Ala Tyr Asn Gly Asn Thr Ala Leu His Val Ala Ala Ser Leu Gln 385 390 395 400

Tyr Arg Leu Thr Gln Leu Asp Ala Val Arg Leu Leu Met Arg Lys Gly 405 410 415

Ala Asp Pro Ser Thr Arg Asn Leu Glu Asn Glu Gln Pro Val His Leu 420 425 430

Val Pro Asp Gly Pro Val Gly Glu Gln Ile Arg Arg Ile Leu Lys Gly

277

435 440 445

Lys Ser Ile Gln Gln Arg Ala Pro Pro Tyr 450 455

<210> 206

<211> 439

<212> PRT

<213> Homo sapien

<400> 206

Trp Ile Val Val Ala Ala Arg Tyr Gln Tyr Ser Pro Gln Asn Gln His 1 5 10 15

Val Glu Gln Gln Pro His Tyr Thr His Lys Pro Thr Leu Glu Tyr Ser 20 25 30

Pro Phe Pro Ile Pro Pro Gln Ser Pro Ala Tyr Glu Pro Asn Leu Phe 35 40 45

Asp Gly Pro Glu Ser Gln Phe Cys Pro Asn Gln Ser Leu Val Ser Leu 50 55 60

Leu Gly Asp Gln Arg Glu Ser Glu Asn Ile Ala Asn Pro Met Gln Thr 65 70 . 75 80

Ser Ser Ser Val Gln Gln Gln Asn Asp Ala His Leu His Ser Phe Ser 85 90 95

Met Met Pro Ser Ser Ala Cys Glu Ala Met Val Gly His Glu Met Ala 100 105 110

Ser Asp Ser Ser Asn Thr Ser Leu Pro Phe Ser Asn Met Gly Asn Pro 115 120 125

Met Asn Thr Thr Gln Leu Gly Lys Ser Leu Phe Gln Trp Gln Val Glu 130 135 140

Gln Glu Glu Ser Lys Leu Ala Asn Ile Ser Gln Asp Gln Phe Leu Ser 145 150 155 160

Lys Asp Ala Asp Gly Asp Thr Phe Leu His Ile Ala Val Ala Gln Gly 165 170 175

Arg Arg Ala Leu Ser Tyr Val Leu Ala Arg Lys Met Asn Ala Leu His 180 185 190

Met Leu Asp Ile Lys Glu His Asn Gly Gln Ser Ala Phe Gln Val Ala 195 200 205

Val Ala Ala Asn Gln His Leu Ile Val Gln Asp Leu Val Asn Ile Gly 210 215 220

Ala Gln Val Asn Thr Thr Asp Cys Trp Gly Arg Thr Pro Leu His Val 225 230 235 240

Cys Ala Glu Lys Gly His Ser Gln Val Leu Gln Ala Ile Gln Lys Gly 245 250 255

Ala Val Gly Ser Asn Gln Phe Val Asp Leu Glu Ala Thr Asn Tyr Asp 260 255 270

Gly Leu Thr Pro Leu His Cys Ala Val Ile Ala His Asn Ala Val Val 275 280 285

His Glu Leu Gln Arg Asn Gln Gln Pro His Ser Pro Glu Val Gln Glu 290 295 300

Leu Leu Leu Lys Asn Lys Ser Leu Val Asp Thr Ile Lys Cys Leu Ile 305 310 315

Gln Met Gly Ala Ala Val Glu Ala Lys Asp Arg Lys Ser Gly Arg Thr 325 330 335

Ala Leu His Leu Ala Ala Glu Glu Ala Asn Leu Glu Leu Ile Arg Leu 340 345 350

Phe Leu Glu Leu Pro Ser Cys Leu Ser Phe Val Asn Ala Lys Ala Tyr 355 360 365

Asn Gly Asn Thr Ala Leu His Val Ala Ala Ser Leu Gln Tyr Arg Leu 370 380

Thr Gln Leu Asp Ala Val Arg Leu Leu Met Arg Lys Gly Ala Asp Pro 385 390 395

Ser Thr Arg Asn Leu Glu Asn Glu Gln Pro Val His Leu Val Pro Asp 405 410 , 415

Gly Pro Val Gly Glu Gln Ile Arg Arg Ile Leu Lys Gly Lys Ser Ile 420 425 430

Gln Gln Arg Ala Pro Pro Tyr 435

<210> 207

<211> 130

<212> PRT

<213> Homo sapien

<400> 207

Met Gln Pro Leu Trp Leu Cys Trp Ala Leu Trp Val Leu Pro Leu Ala

Ser Pro Gly Ala Ala Leu Thr Gly Glu Gln Leu Leu Gly Ser Leu Leu 25

Arg Gln Leu Gln Leu Lys Glu Val Pro Thr Leu Asp Arg Ala Asp Met

Glu Glu Leu Val Ile Pro Thr His Val Arg Ala Gln Tyr Val Ala Leu

Leu Gln Arg Ser His Gly Asp Arg Ser Arg Gly Lys Arg Phe Ser Gln 70

Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Leu Glu Ala Ser Thr

His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro Asn Ser Glu

Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Cys Thr Leu Thr Cys 120

Arg Gly 130

<210> 208 <211> 243 <212> PRT <213> Homo sapien

<400> 208

Asp Pro Pro Ala Ala Phe Ser Arg Asp Ser Pro Thr Leu Pro Leu Ala 1 5 10

Pro Pro Gly Gln His His Ala Ala Pro Val Ala Leu Leu Gly Thr Leu 20 25

| Gly | Val | Ala | Pro | Gly | Gln | Pro | Arg | Gly | Arg | Pro | Asp | Arg | Gly | Ala | Ala |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |

Pro Gly Gln Pro Ala Ala Ala Ala Ala Gln Arg Gly Ala His Pro 50 55 60

Gly Gln Gly Arg His Gly Gly Ala Gly His Pro His Pro Arg Glu Gly 65 70 75 80

Pro Val Arg Gly Pro Ala Ala Ala Gln Pro Arg Gly Pro Leu Pro Arg 85 90 95

Lys Glu Val Gln Pro Glu Leu Pro Arg Gly Gly Arg Gln Val Pro Gly 100 \$105\$

Val Gly Gln His Thr Pro Ala Gly Val Arg His Gly Ala Ala Ala 115 120 125

Ala Ala Gln Gln Arg Ala Gly Ala Gly Arg Ala Ala Leu Pro Gly 130 135 140

Glu Pro Pro Gly Phe Leu Ala Tyr Glu Cys Val Gly Thr Cys Arg Gln 165 170 175

Pro Pro Glu Ala Leu Ala Phe Lys Trp Pro Phe Leu Gly Pro Arg Gln 180 185 190

Cys Ile Ala Ser Glu Thr Asp Ser Leu Pro Met Ile Val Ser Ile Lys 195 200 205

Glu Gly Gly Arg Thr Arg Pro Gln Val Val Ser Leu Pro Asn Met Arg 210 215 220

Val Gln Lys Cys Ser Cys Ala Ser Asp Gly Ala Leu Val Pro Arg Arg 225 230 235 240

Leu Gln Pro

<210> 209

<211> 204

<212> PRT

<213> Homo sapien

<400> 209

Met Glu Arg Leu Thr Leu Pro Leu Gly Gly Ala Ala Ala Val Asp Glu 1 5 10 15

Tyr Leu Glu Tyr Arg Arg Tyr Lys Gln His Lys Thr Asp Leu Glu Ala 20 25 30

Ile Pro Gln Gln Cys Pro Ile Asp Leu Pro Cys Gln Val Thr Gly Cys 35 40 45

Gln Cys Arg Ala Tyr Leu Tyr Val Pro Leu Asn Gly Ser Gln Pro Ile 50 55 60

Arg Cys Arg Cys Lys His Phe Ala Asp Gln His Ser Ala Ala Pro Gly 65 70 75 80

Phe Thr Cys Asn Thr Cys Ser Lys Cys Ser Gly Phe His Ser Cys Phe 85 90 95

Thr Cys Ala Cys Gly Gln Pro Ala Tyr Ala His Asp Thr Val Val Glu 100 105 110

Thr Lys Gln Glu Arg Leu Ala Gln Glu Lys Pro Val Gly Gln Asp Ile 115 120 125

Pro Tyr Ala Ala Met Gly Gly Leu Thr Gly Phe Ser Ser Leu Ala Glu 130 135 140

Gly Tyr Met Arg Leu Asp Asp Ser Gly Ile Val Gly Thr Ser Ser Gln 145 150 155 160

Val Ser Ser Leu Arg Arg Pro Glu Glu Asp Asp Met Ala Phe Phe Glu 165 170 175

Arg Arg Tyr Gln Glu Arg Met Lys Met Glu Lys Ala Ala Lys Trp Lys 180 185 190

Gly Lys Ala Pro Leu Pro Ser Ala Thr Lys Pro Ser

<210> 210

<211> 80

<212> PRT

<213> Homo sapien

<400> 210

282

Glu Val Gln Glu Ala Ile Phe Phe Arg Val Cys Gly Ala Arg Ser Val 1 5 10 15

Val Leu Leu Val Ala Val Arg Leu His Thr Leu Leu Ser Cys Pro 20 25 30

Leu Glu Gln Pro Ala Gly Thr Glu Trp Ile Leu Glu Glu Gly Val Thr 35 40 45

Thr Gly Pro Pro Arg Lys Pro Arg Ala Asp Ile Tyr Asn Leu Arg Ser 50 55 60

Pro Asp Glu Phe Ile Val Gly Gln Asn Gln Ala Leu Ile Glu Pro Gly 65 70 75 80

<210> 211

=

<211> 84

<212> PRT

<213> Homo sapien

<400> 211

Glu Gln Gln Pro Ser Pro Ile Asp Ser Thr Glu Thr Thr Arg Asn Gln
1 5 10 15

Gln Val Arg Pro Thr Thr Ser Arg Asn Lys Arg Arg Ala Ala Ser Gln 20 25 30

His Ile Ser Lys Ala Thr Arg Pro Thr Ala Lys Arg Gln Ala Ala Asp

Ser Asp Ile Thr Thr Ser Gly Pro Thr Ala Thr Thr Thr Asp Asp Lys 50 55 60

Asn Asp Val Cys Glu Asp Thr Pro His Arg Arg Thr Thr Gly Trp His 65 70 75 80

Gln Arg Asp Leu

<210> 212

<211> 56

<212> PRT

<213> Homo sapien

<400> 212

Pro Leu Trp Arg Arg Leu Leu Leu Gly Ser Arg Leu Leu Leu Pro Cys
1 5 10 15

Asn Arg Asn Trp Arg Trp Asn Met Arg Gly Ala Leu Trp Lys Glu Lys 20 25

Asp Arg Pro Cys Ala Phe Met Lys Val Lys Ile Trp Leu Asn Gln Phe 35

His Lys Val Thr Val Tyr Ile Ala 50

<210> 213

<211> 451

<212> PRT

<213> Homo sapien

<400> 213

Met Phe Leu Leu Leu His Leu Gln Ile Lys Trp Arg Ala Thr Ile 5

Asn Leu Leu Ser Val Thr Glu Asp Gly Leu His Phe Val Glu Tyr Tyr 25

Leu Asn Arg Ile Ile His Leu Asp Val Asp Ser Glu Ala Lys Lys Leu 40

Leu Gly Leu Gly Gln Lys His Leu Val Met Gly Asp Ile Pro Ala Ala

Val Asn Ala Phe Gln Glu Ala Ala Ser Leu Leu Gly Lys Lys Tyr Gly 70 75

Glu Thr Ala Asn Glu Cys Gly Glu Ala Phe Phe Phe Tyr Gly Lys Ser 90

Leu Leu Glu Leu Ala Arg Met Glu Asn Gly Val Leu Gly Asn Ala Leu 105 100

Glu Gly Val His Val Glu Glu Glu Glu Gly Glu Lys Thr Glu Asp Glu 115

Ser Leu Val Glu Asn Asn Asp Asn Ile Asp Glu Thr Glu Gly Ser Glu 130 135

Glu Asp Asp Lys Glu Asn Asp Lys Thr Glu Glu Met Pro Asn Asp Ser 150

284

Val Leu Glu Asn Lys Ser Leu Gln Glu Asn Glu Glu Glu Glu Ile Gly 165 170 175

Asn Leu Glu Leu Ala Trp Asp Met Leu Asp Leu Ala Lys Ile Ile Phe 180 185 190

Lys Arg Gln Glu Thr Lys Glu Ala Gln Leu Tyr Ala Ala Gln Ala His 195 200 205

Leu Lys Leu Gly Glu Val Ser Val Glu Ser Glu Asn Tyr Val Gln Ala 210 215 220

Val Glu Glu Phe Gln Ser Cys Leu Asn Leu Gln Glu Gln Tyr Leu Glu 225 235 240

Ala His Asp Arg Leu Leu Ala Glu Thr His Tyr Gln Leu Gly Leu Ala 245 250 255

Tyr Gly Tyr Asn Ser Gln Tyr Asp Glu Ala Val Ala Gln Phe Ser Lys 260 265 270

Ser Ile Glu Val Ile Glu Asn Arg Met Ala Val Leu Asn Glu Gln Val 275 280 285

Lys Glu Ala Glu Gly Ser Ser Ala Glu Tyr Lys Lys Glu Ile Glu Glu 290 295 300

Leu Lys Glu Leu Leu Pro Glu Ile Arg Glu Lys Ile Glu Asp Ala Lys 305 310 315

Glu Ser Gln Arg Ser Gly Asn Val Ala Glu Leu Ala Leu Lys Ala Thr 325 330 335

Leu Val Glu Ser Ser Thr Ser Gly Fhe Thr Pro Gly Gly Gly Ser 340 345 350

Ser Val Ser Met Ile Ala Ser Arg Lys Pro Thr Asp Gly Ala Ser Ser 355 360 365

Ser Asn Cys Val Thr Asp Ile Ser His Leu Val Arg Lys Lys Arg Lys 370 375 380

Pro Glu Glu Glu Ser Pro Arg Lys Asp Asp Ala Lys Lys Ala Lys Gin 385 390 395 400

Glu Pro Glu Val Asn Gly Gly Ser Gly Asp Ala Val Pro Ser Gly Asn

415 410 405

Glu Val Ser Glu Asn Met Glu Glu Glu Ala Glu Asn Gln Ala Glu Ser 420 425

Arg Ala Ala Val Glu Gly Thr Val Glu Ala Gly Ala Thr Val Glu Ser 440

Thr Ala Cys 450

<210> 214

<211> 214 <211> 337 <212> PRT <213> Homo sapien

<400> 214

Met Ala His Ala Pro Ala Arg Cys Pro Ser Ala Arg Gly Ser Gly Asp

Gly Glu Met Gly Lys Pro Arg Asn Val Ala Leu Ile Thr Gly Ile Thr 20

Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr

Glu Val His Gly Ile Val Arg Arg Ser Ser Phe Asn Thr Gly Arg 55

Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met 70

Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile 85 90

Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser 100 . 105 110

His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp 115 120 125

Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Val Lys Thr Cys Gly Leu 130 135 140

Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly 155 160

Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg 165 170

Ser Pro Tyr Gly Ala Asn Phe Val Thr Arg Lys Ile Ser Arg Ser Val 180 185

Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu Gly Asn Leu

Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val Glu Ala Met 210 215

Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val Ile Ala Thr 225 230 235 240

Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser Phe Leu His 245 250 255

Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn Glu Val Gly 260

Arg Cys Lys Glu Thr Gly Lys Val His Val Thr Val Asp Leu Lys Tyr 275 280

Tyr Arg Pro Thr Glu Val Asp Phe Leu Gln Gly Asp Cys Thr Lys Ala 290 295

Lys Gln Lys Leu Asn Trp Lys Pro Arg Val Ala Phe Asp Glu Leu Val 305 310

Arg Glu Met Val His Ala Asp Val Glu Leu Met Arg Thr Asn Pro Asn 325

Ala

<210> 215

<211> 332

<211> 332
<212> PRT
<213> Homo sapien

<400> 215

Met Ala His Ala Pro Ala Arg Cys Pro Ser Ala Arg Gly Ser Gly Asp

Gly Glu Met Gly Lys Pro Arg Asn Val Ala Leu Ile Thr Gly Ile Thr

20 25 30

Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr 35 40 45

Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg 50 60

Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met 65 70 75 80

Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile 85 90 95

Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser 100 105 110

His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp 115 120 125

Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Val Lys Thr Cys Gly Leu 130 135 140

Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly 145 150 155

Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg 165 170 175

Ser Pro Tyr Gly Ala Ala Lys Leu Tyr Ala Tyr Trp Ile Val Val Asn 180 185 190

Phe Arg Glu Ala Tyr Asn Leu Phe Ala Val Asn Gly Ile Leu Phe Asn 195 200 205

His Glu Ser Pro Arg Arg Gly Ala Asn Phe Val Thr Arg Lys Ile Ser 210 215 220

Arg Ser Val Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu 225 230 235 240

Gly Asn Leu Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val 245 250 255

Glu Ala Met Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val 260 265 270

Ile Ala Thr Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser 275 280 285

Phe Leu His Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn 290 295 300

Glu Val Gly Arg Cys Lys Glu Thr Gly Lys Val His Val Thr Val Asp 305 310 315 320

Leu Lys Tyr Tyr Arg Pro Thr Glu Val Glu Thr Asn 325

<210> 216

<211> 382

<212> PRT

<213> Homo sapien

<400> 216

Met Ala His Ala Pro Ala Arg Cys Pro Ser Ala Arg Gly Ser Gly Asp 1 5 10 15

Gly Glu Met Gly Lys Pro Arg Asn Val Ala Leu Ile Thr Gly Ile Thr 20 25 30

Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr 35 40 45

Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg 50 55 60

Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met 65 70 75 80

Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile 85 90 95

Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser 100 105 110

His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp

Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Val Lys Thr Cys Gly Leu 130 135 140

Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly 150 Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg Ser Pro Tyr Gly Ala Ala Lys Leu Tyr Ala Tyr Trp Ile Val Val Asn 180 185 190 Phe Arg Glu Ala Tyr Asn Leu Phe Ala Val Asn Gly Ile Leu Phe Asn 195 200 His Glu Ser Pro Arg Arg Gly Ala Asn Phe Val Thr Arg Lys Ile Ser 210 215 220 Arg Ser Val Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu 235 230 225 Gly Asn Leu Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val 245 250 255 Glu Ala Met Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val 260 265 Ile Ala Thr Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser 275 Phe Leu His Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn 290 295 300 Glu Val Gly Arg Cys Lys Glu Thr Gly Lys Val His Val Thr Val Asp 305 310 Leu Lys Tyr Tyr Arg Pro Thr Glu Val Val Arg Thr Leu Trp Pro Pro 325 335 Ser Ala Trp Pro Arg Leu Ala Gly Trp Leu Gly Lys Cys Ala His Gly 345 350 340

Met Pro Gly Ala Ser Leu Trp Ser Cys Gln Phe Ser Ser Leu Ala Ser

365

Phe Ser Val His Phe Gln Asn Gln Asn Thr Val Asn Ser Ile 370 375 380

360

355

<211> 258

<212> PRT

<213> Homo sapien

<400> 217

Met Asn Ser Asn Val Glu Asn Leu Pro Pro His Ile Ile Arg Leu Val 1 5 10 15

Tyr Lys Glu Val Thr Thr Leu Thr Ala Asp Pro Pro Asp Gly Ile Lys 20 25 30

Val Phe Pro Asn Glu Glu Asp Leu Thr Asp Leu Gln Val Thr Ile Glu 35 40 45

Gly Pro Glu Gly Thr Pro Tyr Ala Gly Gly Leu Phe Arg Met Lys Leu  $50 \hspace{1cm} 55 \hspace{1cm} 60$ 

Leu Leu Gly Lys Asp Phe Pro Ala Ser Pro Pro Lys Gly Tyr Phe Leu 65 70 75 80

Thr Lys Ile Phe His Pro Asn Val Gly Ala Asn Gly Glu Ile Cys Val 85 90 95

Asn Val Leu Lys Arg Asp Trp Thr Ala Glu Leu Gly Ile Arg His Val

Leu Leu Thr Ile Lys Cys Leu Leu Ile His Pro Asn Pro Glu Ser Ala 115 120 125

Leu Asn Glu Glu Ala Gly Arg Leu Leu Glu Asn Tyr Glu Glu Tyr 130 135 140

Ala Ala Arg Ala Arg Leu Leu Thr Glu Ile His Gly Gly Ala Gly Gly 145 150 155 160

Pro Ser Gly Arg Ala Glu Ala Gly Arg Ala Leu Ala Ser Gly Thr Glu 165 170 175

Ala Ser Ser Thr Asp Pro Gly Ala Pro Gly Arg Arg Ala Glu Val His 180 . 185 190

Trp Pro Glu His Val Gly Arg Glu Arg Arg Trp Gly Arg Lys Thr Thr
195 200 205

Asp Gly Ala Arg Val Lys Val Phe Leu Ser Arg Asp His Ser Ala Pro 210 215 220

Asn Phe Ser Asn Cys Gly Pro Ser Gly Arg Arg Val-Asn Ala Gln Thr 225 230 235 240

Lys Lys Pro Ser Arg Lys Gly Val Leu Ser Ala Ala Phe Gln Ala Ser 245 250 255

Leu Leu

<210> 218

<211> 262

<212> PRT

<213> Homo sapien

<400> 218

Met Arg Ala Val Ile Lys Arg Gln Trp Cys Ala Arg Gly Arg Leu Ser 1 5 10 15

Ala Ala Gly His Arg Gly Gly Leu Val Tyr Ala Val Arg Gly Gly 20 25 30

Arg Arg Arg Gln Arg Gly Ala Glu Arg Gly Arg Arg Gly Leu Ser Arg 35 40 45

Ala Ala Ala Ala Val Gly Pro Pro Ala Pro Ala Gly Gly Pro Lys 50 55 60

Asn Leu Pro Pro His Ile Ile Arg Leu Val Tyr Lys Glu Val Thr Thr 65 70 75 80

Leu Thr Ala Asp Pro Pro Asp Gly Ile Lys Val Phe Pro Asn Glu Glu 85 90 95

Asp Leu Thr Asp Leu Gln Val Thr Ile Glu Gly Pro Glu Gly Thr Pro 100 105 110

Tyr Ala Gly Gly Leu Phe Arg Met Lys Leu Leu Gly Lys Asp Phe 115 120 125

Pro Ala Ser Pro Pro Lys Gly Tyr Phe Leu Thr Lys Ile Phe His Pro 130 135 140

Asn Val Gly Ala Asn Gly Glu Ile Cys Val Asn Val Leu Lys Arg Asp 145 150 155 160

Trp Thr Ala Glu Leu Gly Ile Arg His Val Leu Leu Thr Ile Lys Cys

|  |  | • |
|--|--|---|
|  |  |   |

175 165 170

Leu Leu Ile His Pro Asn Pro Glu Ser Ala Leu Asn Glu Glu Ala Gly 185 180

Arg Leu Leu Glu Asn Tyr Glu Glu Tyr Ala Ala Arg Ala Arg Leu 195 200 205

Leu Thr Glu Ile His Gly Gly Ala Gly Gly Pro Ser Gly Arg Ala Glu 210 215 220

Ala Gly Arg Ala Leu Ala Ser Gly Thr Glu Ala Ser Ser Thr Asp Pro 230 235 240

Gly Ala Pro Gly Gly Pro Gly Gly Ala Glu Gly Ser His Gly Gln Glu 245 250

Ala Cys Trp Arg Ala Arg 260

<210> 219 <211> 291 <212> PRT <213> Homo sapien

<400> 219

Gly Ser Glu Leu Arg Gly Arg Gly Arg Gly Leu Arg Ala Val Ile

Lys Arg Gln Trp Cys Ala Arg Gly Arg Leu Ser Ala Ala Gly His Arg 20 25

Gly Gly Gly Leu Val Tyr Ala Val Arg Gly Gly Arg Arg Arg Gln Arg 40

Gly Ala Glu Arg Gly Arg Gly Leu Ser Arg Ala Ala Ala Ala Ala 55

Val Gly Pro Pro Ala Pro Ala Gly Gly Pro Lys Asn Leu Pro Pro His 70 ' 75

Ile Ile Arg Leu Val Tyr Lys Glu Val Thr Thr Leu Thr Ala Asp Pro 85 90

Pro Asp Gly Ile Lys Val Phe Pro Asn Glu Glu Asp Leu Thr Asp Leu 105

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Gln Val Thr Ile Glu Gly Pro Glu Gly Thr Pro Tyr Ala Gly Gly Leu 120

Phe Arg Met Lys Leu Leu Cly Lys Asp Phe Pro Ala Ser Pro Pro

Lys Gly Tyr Phe Leu Thr Lys Ile Phe His Pro Asn Val Gly Ala Asn

Gly Glu Ile Cys Val Asn Val Leu Lys Arg Asp Trp Thr Ala Glu Leu

Gly Ile Arg His Val Leu Leu Thr Ile Lys Cys Leu Leu Ile His Pro 180 185

Asn Pro Glu Ser Ala Leu Asn Glu Glu Ala Gly Arg Leu Leu Glu 195 200 205

Asn Tyr Glu Glu Tyr Ala Ala Arg Ala Arg Leu Leu Thr Glu Ile His 215

Gly Gly Ala Gly Gly Pro Ser Gly Arg Ala Glu Ala Gly Arg Ala Leu

Ala Ser Gly Thr Glu Ala Ser Ser Thr Asp Pro Gly Ala Pro Gly Gly 250

Pro Gly Gly Ala Glu Gly Pro Met Ala Lys Lys His Ala Gly Glu Arg

Asp Lys Lys Leu Ala Ala Lys Lys Lys Thr Asp Lys Lys Arg Ala Leu

Arg Arg Leu 290

<210> 220

<211> 233 <212> PRT

<213> Homo sapien

<400> 220

Met Asn Ser Asn Val Glu Asn Leu Pro Pro His Ile Ile Arg Leu Val 5

Tyr Lys Glu Val Thr Thr Leu Thr Ala Asp Pro Pro Asp Gly Ile Lys

20

25

30

Val Phe Pro Asn Glu Glu Asp Leu Thr Asp Leu Gln Val Thr Ile Glu 35 40 45

Gly Pro Glu Gly Thr Pro Tyr Ala Gly Gly Leu Phe Arg Met Lys Leu 50 55 60

Leu Leu Gly Lys Asp Phe Pro Ala Ser Pro Pro Lys Gly Tyr Phe Leu 65 70 75 80

Thr Lys Ile Phe His Pro Asn Val Gly Ala Asn Gly Glu Ile Cys Val 85 90 95

Asn Val Leu Lys Arg Asp Trp Thr Ala Glu Leu Gly Ile Arg His Val

Leu Leu Ser Trp Lys Asp Lys Gln Cys Gln Thr Gln Asp Thr Gln
115 120 125

Val Leu Leu Arg Ser Ala Gln Glu His Leu Thr Met Gln Arg Val Thr 130 140

Ile Lys Cys Leu Leu Ile His Pro Asn Pro Glu Ser Ala Leu Asn Glu 145 150 155 160

Glu Ala Gly Arg Leu Leu Glu Asn Tyr Glu Glu Tyr Ala Ala Arg 165 170 175

Ala Arg Leu Eu Thr Glu Ile His Gly Gly Ala Gly Gly Pro Ser Gly 180 135 190

Arg Ala Glu Ala Gly Arg Ala Leu Ala Ser Gly Thr Glu Ala Ser Ser 195 200 205

Thr Asp Pro Gly Ala Pro Gly Gly Pro Gly Gly Ala Glu Gly Ser His 210 220

Gly Gln Glu Ala Cys Trp Arg Ala Arg 225 230

<210> 221

<211> 390

<212> PRT

<213> Homo sapien

<400> 221

Glu Pro Ser Arg Pro Pro Arg Ala Pro Ile Gly Arg Pro Ala Thr Gln

Pro Ser Pro Pro Leu Pro Ser Leu Leu Thr Ile Val Cys Gly Glu Gly

Ser Glu Arg Val Glu Asn Gln Gly Thr Cys Ala Leu Thr Ser Arg Leu

Arg Leu Gly Ala Glu Gly Pro Arg Arg Ala Trp Pro Ala Gly Gly Tyr

Lys Glu Ala Val Val Arg Ala Arg Pro Ala Gln Cys Cys Arg Ala Pro

Gly Arg Arg Val Gly Leu Arg Cys Ala Arg Arg Thr Ser Glu Ala Ala

Gly Ser Gly Ala Gly Pro Pro Gly Pro Leu Gln Gly Arg Ser Gly Ser 100 105

Ser Trp Ala Pro Arg Pro Gly Arg Arg Thr Glu Glu Arg Arg Lys Gly 120 115

Ala Gly Gly Thr Arg Pro Arg Pro Ala Ala Ala Met Asn Ser Asn Val 135 130

Glu Asn Leu Pro Pro His Ile Ile Arg Leu Val Tyr Lys Glu Val Thr 155 145

Thr Leu Thr Ala Asp Pro Pro Asp Gly Ile Lys Val Phe Pro Asn Glu 170

Glu Asp Leu Thr Asp Leu Gln Val Thr Ile Glu Gly Pro Glu Gly Thr 185

Pro Tyr Ala Gly Gly Leu Phe Arg Met Lys Leu Leu Gly Lys Asp

Phe Pro Ala Ser Pro Pro Lys Gly Tyr Phe Leu Thr Lys Ile Phe His 215 . 220

Pro Asn Val Gly Ala Asn Gly Glu Ile Cys Val Asn Val Leu Lys Arg 230 235

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Asp Trp Thr Ala Glu Leu Gly Ile Arg His Val Leu Leu Leu Ser Trp 245 250 255

Lys Asp Lys Gln Cys Gln Thr Gln Asp Thr Gln Val Leu Leu Arg Ser 260 265 270

Ala Gln Glu His Leu Thr Met Gln Arg Val Thr Ile Lys Cys Leu Leu 275 280 285

Ile His Pro Asn Pro Glu Ser Ala Leu Asn Glu Glu Ala Gly Arg Leu 290 295 300

Leu Leu Glu Asn Tyr Glu Glu Tyr Ala Ala Arg Ala Arg Leu Leu Thr 305 310 315

Arg Ala Leu Ala Ser Gly Thr Glu Ala Ser Ser Thr Asp Pro Gly Ala 340 345 350

Pro Gly Gly Pro Gly Gly Ala Glu Gly Pro Met Ala Lys Lys His Ala 355 360 365

Gly Glu Arg Asp Lys Lys Leu Ala Ala Lys Lys Lys Thr Asp Lys Lys 370 \$375\$

Arg Ala Leu Arg Arg Leu 385 390

<210> 222

<211> 110

<212> PRT

<213> Homo sapien

<400> 222

Pro Gly Ala His Pro Ser Leu Asp Leu Thr Arg Cys Ser Leu Phe Leu 1 10 15

Pro Lys Arg Ala Arg Ser Ala Ile Thr Lys Ile Ser Leu Val Leu Tyr
20 25 30

Phe Leu Thr Arg Arg Arg Thr Gly Trp Arg Cys Ser Pro Ala Ala 35 40 45

Trp Arg Cys Gln Arg Ser Glu Gly Leu Gln Glu Gly Leu Lys Leu Pro
50 55 60

Ala Gln Asn Leu Arg Met Glu Pro Ala Leu His Tyr Leu Arg Ser Gln 70 75

Gly Leu Gly Arg Trp Arg Lys Val Ile Ser Pro Ser Leu Lys Ser Tyr 85 90

Phe Leu Asn Val Ala Pro His Gln Ala Leu Tyr Leu Thr Ser 105 110

<210> 223 <211> 257

<212> PRT

<213> Homo sapien

<400> 223

Met Asp His Arg Ser Arg Leu Arg Gly Thr Gly Leu Asn Arg Ile Pro 10

Gly Thr Gln Ser Arg Ala Pro Arg Val Pro Leu Pro Phe His Val Gln 25

Gln Glu Ala Arg Glu Gly Glu Asp Trp Glu Arg Glu Pro Pro Arg Gln

Arg Pro Pro Ile Tyr Glu Pro Pro Glu Ser Glu Glu Leu Pro Asp Asn

Val Met Val Ser Lys Pro Ala Pro Tyr Trp Glu Gly Thr Ala Val Ile

Asp Gly Glu Phe Lys Glu Leu Lys Leu Thr Asp Tyr Arg Gly Lys Tyr

Leu Val Phe Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro Thr 100 105

Glu Ile Ile Ala Phe Gly Asp Arg Leu Glu Glu Phe Arg Ser Ile Asn 115 120

Thr Glu Val Val Ala Cys Ser Val Asp Ser Gln Phe Thr His Leu Ala 140

Trp Ile Asn Thr Pro Arg Arg Gln Gly Gly Leu Gly Pro Ile Arg Ile 145 150 155

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Pro Leu Leu Ser Asp Leu Thr His Gln Ile Ser Lys Asp Tyr Gly Val 170

Tyr Leu Glu Asp Ser Gly His Thr Leu Arg Gly Leu Phe Ile Ile Asp 185

Asp Lys Gly Ile Leu Arg Gln Ile Thr Leu Asn Asp Leu Pro Val Gly 195 200

Arg Ser Val Asp Glu Thr Leu Arg Leu Val Gln Ala Phe Gln Tyr Thr 215

Asp Lys His Gly Glu Val Cys Pro Ala Gly Trp Lys Pro Gly Ser Glu

Thr Ile Ile Pro Asp Pro Ala Gly Lys Leu Lys Tyr Phe Asp Lys Leu 250

Asn

<210> 224 <211> 105 <212> PRT <213> Homo sapien

<400> 224

Met Gln Lys Lys Lys Asn Ser Asn Ser Asn Ser Gly Thr Ser Ser Phe

Gly Lys Arg Arg Asn Lys Thr His Thr Leu Cys Arg Arg Cys Gly Ser 20

Lys Ala Tyr His Leu Gln Lys Ser Thr Cys Gly Lys Cys Gly Tyr Pro 40

Ala Lys Arg Lys Arg Lys Tyr Asn Trp Ser Ala Lys Ala Lys Arg Arg

Asn Thr Thr Gly Thr Gly Arg Met Arg His Leu Lys Ile Val Tyr Arg

Arg Phe Arg His Gly Phe Arg Glu Gly Thr Thr Pro Lys Pro Lys Arg 90

> Ala Ala Val Ala Ala Ser Ser Ser 100

<210> 225 <211> 111 <212> PRT <213> Homo sapien <400> 225 Ile Phe Met Val Gly Val Asp Ala Lys Lys Lys Glu Phe Glu Phe Glu Phe Gly Thr Ser Ser Phe Gly Lys Arg Arg Asn Lys Thr His Thr Leu Cys Arg Arg Cys Gly Ser Lys Ala Tyr His Leu Gln Lys Ser Thr Cys 40 Gly Lys Cys Gly Tyr Pro Ala Lys Arg Lys Arg Lys Tyr Asn Trp Ser Ala Lys Ala Lys Arg Arg Asn Thr Thr Gly Thr Gly Arg Met Arg His 70 Leu Lys Ile Val Tyr Arg Arg Phe Arg His Gly Phe Arg Glu Gly Thr 90 Thr Pro Lys Pro Lys Arg Ala Ala Val Ala Ala Ser Ser Ser Ser 105 <210> 226 <211> 104 <212> PRT <213> Homo sapien <220> <221> MISC\_FEATURE <222> (3)..(5) <223> X=any amino acid <400> 226

Met Ser Xaa Xaa Xaa Arg Ile Arg Pro Arg Gly Thr Ser Ser Phe Gly 1 5 10 15

Lys Arg Arg Asn Lys Thr His Thr Leu Cys Arg Arg Cys Gly Ser Lys 20 25 30

Ala Tyr His Leu Gln Lys Ser Thr Cys Gly Lys Cys Gly Tyr Pro Ala 35 40 45

Lys Arg Lys Arg Lys Tyr Asn Trp Ser Ala Lys Ala Lys Arg Arg Asn 55

Thr Thr Gly Thr Gly Arg Met Arg His Leu Lys Ile Val Tyr Arg Arg 70

Phe Arg His Gly Phe Arg Glu Gly Thr Thr Pro Lys Pro Lys Arg Ala 95

Ala Val Ala Ala Ser Ser Ser Ser 100

<210> 227

<211> 129

<212> PRT

<213> Homo sapien

<220>

<221> MISC\_FEATURE

<222> (12)..(12)

<223> X=any amino acid

<220>

<221> MISC\_FEATURE

<222> (25)..(25)

<223> X=any amino acid

<220>

<221> MISC\_FEATURE

<222> (62)..(62) <223> X=any amino acid

<220>

<221> MISC\_FEATURE

<222> (64)..(64) <223> X=any amino acid

<400> 227

Gln Ser His Lys Thr Leu Val Leu Gln Thr Thr Xaa Arg Ser Leu Leu 5

Ala His Thr Thr Cys Arg Phe Trp Xaa Phe Pro Asn Leu Leu Gly Ile 20 25

Lys Val Asn Asn Ser Ile Thr Arg Gly Ser Gly Gln Pro Ser Phe Val 40

Arg Gly Cys Ile Val Gly Lys Pro Thr Ser Val Cys Gln Xaa Leu Xaa

60 55 50

Glu Phe Gly Arg Gly Glu Arg His Arg Leu Glu Ser Val Ala Ile Arg 70

Arg Thr Arg Cys Ala Ala Ala Val Ala Leu Arg Pro Thr Thr Phe Arg

Ser Arg Pro Val Ala Asn Val Ala Thr Leu Pro Ser Ala Arg Glu Ser

Ile Thr Gly Val Pro Arg Leu Lys Asp Glu Ile Pro Pro Glu Leu Val 120

Glu

<210> 228

<211> 96

<212> PRT

<213> Homo sapien

<400> 228

Ala Cys Arg Ala Ala Gln Cys Asp Gly Ser Trp Ser Arg Pro Arg Ser 5 10

Leu Cys Arg Arg Cys Gly Ser Lys Ala Tyr His Leu Gln Lys Ser Thr 25 20

Cys Gly Lys Cys Gly Tyr Pro Ala Lys Arg Lys Arg Lys Tyr Asn Trp 35 40

Ser Ala Lys Ala Lys Arg Arg Asn Thr Thr Gly Thr Gly Arg Met Arg 50 55

His Leu Lys Ile Val Tyr Arg Arg Phe Arg His Gly Phe Arg Glu Gly 70

Thr Thr Pro Lys Pro Lys Arg Ala Ala Val Ala Ala Ser Ser Ser 85

<210> 229 <211> 55 <212> PRT <213> Homo sapien

<400> 229

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302

Met His Ala Glu Arg Arg Ser Val Met Asp Arg Gly Arg Gly Arg Gly 1 5 10 15

Arg Pro Thr Thr Phe Arg Ser Arg Pro Val Ala Asn Val Ala Thr Leu 20 25 30

Pro Ser Ala Arg Glu Ser Ile Thr Gly Val Pro Arg Leu Lys Asp Glu 35 40

Ile Pro Pro Glu Leu Val Glu 50 55

<210> 230

<211> 72

<212> PRT

<213> Homo sapien

<400> 230

Ala Tyr His Leu Gln Lys Ser Thr Cys Gly Lys Cys Gly Tyr Pro Ala 1 5 10 15

Lys Arg Lys Arg Lys Tyr Asn Trp Ser Ala Lys Ala Lys Arg Arg Asn 20 25 30

Thr Thr Gly Thr Gly Arg Met Arg His Leu Lys Ile Val Tyr Arg Arg 35 40 45

Phe Arg His Gly Phe Arg Glu Gly Thr Thr Pro Lys Pro Lys Arg Ala 50 55

Ala Val Ala Ala Ser Ser Ser Ser 65

<210> 231

<211> 185

<212> PRT

<213> Homo sapien

<400> 231

Met Leu Glu Arg Arg Ser Val Asp Gly Cys Ala Arg Ala Gly Gly Arg 1 5 10 15

Ala Gly Gly Ala Ile Met Gly Val Asp Ile Arg His Asn Lys Asp Arg
20 25 30

Lys Val Arg Arg Lys Glu Pro Lys Ser Gln Asp Ile Tyr Leu Arg Leu 35 40 45

Leu Val Lys Leu Tyr Arg Phe Leu Ala Arg Arg Thr Asn Ser Thr Phe

Asn Gln Val Val Leu Lys Arg Leu Phe Met Ser Arg Thr Asn Arg Pro

Pro Leu Ser Leu Ser Arg Met Ile Arg Lys Met Lys Leu Pro Gly Arg 90

Glu Asn Lys Thr Ala Val Val Gly Thr Ile Thr Asp Asp Val Arg 105 100

Val Gln Glu Val Pro Lys Leu Lys Val Cys Ala Leu Arg Val Thr Ser

Arg Ala Arg Ser Arg Ile Leu Arg Ala Gly Gly Lys Ile Leu Thr Phe

Asp Gln Leu Ala Leu Asp Ser Pro Lys Gly Cys Gly Thr Val Leu Leu 150

Ser Gly Pro Arg Lys Gly Arg Glu Val Tyr Arg His Phe Gly Lys Ala 170

Pro Gly Thr Pro His Ser His Thr Lys

<210> 232

<211> 214 <212> PRT

<213> Homo sapien

<400> 232

Gly Leu Trp His Cys Pro Ala Leu Arg Ser Ser Gln Gly Pro Arg Gly 5

Val Pro Ala Phe Arg Gln Gly Pro Arg Asn Pro Ala Gln Pro His Gln 20

Val Ser Ile Arg Pro Pro Ala Leu Pro Ser Pro Gln Thr Gln Pro Ala

Gly Pro Gly Leu Ala Thr Leu Gly Leu Leu Leu Ser Leu Val Pro 50

Ala Ser Pro Arg Pro Ser Gly Thr Leu Ser Cys Leu Ile Leu Pro Ala

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304

75 70 65

Phe Pro Phe Asn Thr Ala Trp Ser Cys Val Phe Gln Gly Leu Ser Arg 90 85

His Leu Leu Gly Ser Met Gln Phe Thr Gly Leu Cys Gln Pro Arg Leu 105 100

Gly Pro Ser Arg Trp Trp Gly Arg Cys Phe His Ser Pro Ser Trp Leu

Leu Gly Phe Pro Leu Cys Gln Ala Phe Pro Ala Ala Leu Thr Leu Leu

Gly Leu Asn Val Thr Gly Leu Trp Cys Ser Cys Ala Thr Pro Gln Trp

Pro Pro Leu Arg Gly Pro Pro Ser His Ser Leu Leu Ser Pro Gln Thr 170

Leu Arg Pro Leu Gln Gly Pro Glu Val Arg Ala Cys Gln Arg Pro Thr

Gly Gln Pro Arg Leu Gln Lys Leu Thr Leu Asp Pro Thr Leu Leu Leu 200

Lys Arg Phe Leu Leu Thr 210

<210> 233 <211> 131 <212> PRT <213> Homo sapien

<400> 233

Met Leu Glu Arg Arg Ser Val Asp Gly Cys Ala Arg Ala Gly Gly Arg 5

Ala Gly Gly Ala Ile Met Gly Val Asp Ile Arg His Asn Lys Asp Arg

Lys Val Arg Arg Lys Glu Pro Lys Ser Gln Asp Ile Tyr Leu Arg Leu 40

Leu Val Lys Leu Tyr Arg Phe Leu Ala Arg Arg Thr Asn Ser Thr Phe 50 55

Asn Gln Val Val Leu Lys Arg Leu Phe Met Ser Arg Thr Asn Arg Pro

Pro Leu Ser Leu Ser Arg Met Ile Arg Lys Met Lys Leu Pro Gly Arg

Glu Asn Lys Thr Ala Val Val Gly Thr Ile Thr Asp Asp Val Arg 100 . 105

Val Gln Glu Val Pro Lys Leu Ile Gly Arg Asp His Ala Lys Pro Asp 120

Ser Ser Thr 130

<210> 234 <211> 132 <212> PRT <213> Homo sapien

<400> 234

Asp Ala Cys Ser Ser Gly Ala Gly Asp Gly Cys Ala Arg Ala Gly Gly 5 10

Arg Ala Gly Gly Ala Ile Met Gly Val Asp Ile Arg His Asn Lys Asp 25

Arg Lys Val Arg Arg Lys Glu Pro Lys Ser Gln Asp Ile Tyr Leu Arg

Leu Leu Val Lys Leu Tyr Arg Phe Leu Ala Arg Arg Thr Asn Ser Thr 50 55

Phe Asn Gln Val Val Leu Lys Arg Leu Phe Met Ser Arg Thr Asn Arg

Pro Pro Leu Ser Leu Ser Arg Met Ile Arg Lys Met Lys Leu Pro Gly

Arg Glu Asn Lys'Thr Ala Val Val Gly Thr Ile Thr Asp Asp Val 100 105

Arg Val Gln Glu Val Pro Lys Leu Ile Gly Arg Asp His Ala Lys Pro 115 120 125

Asp Ser Ser Thr

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130

<210> 235

<211> 195 <212> PRT

<213> Homo sapien

<400> 235

Met Asp Trp Ser Arg Arg Gly Gly Arg Ala Gly Gly Ala Ile Met Gly

Val Asp Ile Arg His Asn Lys Asp Arg Lys Val Arg Arg Lys Glu Pro 25

Lys Ser Gln Asp Ile Tyr Leu Arg Leu Leu Val Lys Leu Tyr Arg Phe

Leu Ala Arg Arg Thr Asn Ser Thr Phe Asn Gln Val Val Leu Lys Arg

Leu Phe Met Ser Arg Thr Asn Arg Pro Pro Leu Ser Leu Ser Arg Met 75 70

Val Ser Gly Trp Ser Arg Glu His Gly Arg Pro Gly Ser Arg Trp Val 85 90

Leu Ser Val Trp Lys Gly Gly Arg Thr Trp Ser Ser Gly Ser Asn Gln 105 100

Gly Ile Lys Gly Leu Ser Gln Pro Val Ala Ser Val Glu Leu Gly Leu 115

Leu Val Gly Thr Glu Cys Pro Trp Ala Val Gly Lys Ser Pro Gly Pro

Pro Leu Leu Phe Val Arg Trp Arg Cys Pro Gly Gly Phe Arg Arg 150 155 160

Leu Pro Gln Val Ile Thr Glu Phe Tyr Val Lys Gly Ser Ala Glu Gly 165 . 170

Gly Pro Ile Glu Gln Ser Ala Phe Phe Phe Leu Ser Gly Ala Phe Pro 180 185 190

Ser Trp Thr 195

<210> 236 <211> 115 <212> PRT <213> Homo sapien

<400> 236

Ser Ala Trp Thr Leu Thr Ser Asn Gly Arg Ser Phe Pro Gly Pro Phe 10 15 5

Pro Lys Ser Trp Ala Cys Phe Leu Leu Pro Leu Ala Ile Leu Cys Pro 20 25

Cys Gly Cys Ser Pro Thr Leu Arg Ala Val Pro Asp Leu Ser Cys Tyr 40 35

Phe Pro Lys Pro Glu Thr Ala Pro Leu Gln Ile Leu Ala Ala Pro Phe 55

Pro Cys Val Gln Tyr Arg Val Ile Thr Cys Pro Ser Leu Val Pro Leu 70 75 80

Ile Leu Pro Cys Asp Tyr Ser Val Ile Pro Leu Pro Val Pro Glu Pro 85 90

Pro Gly Leu Phe Leu Gly Ser Pro Glu Cys Ser Pro Arg Thr Gln Ser

Ala Val Pro 115

<210> 237

<211> 156

<212> PRT

<213> Homo sapien

<400> 237

Gln Ser Leu Gly Arg Gly Leu Ala Thr Thr Arg Gly Arg Gly Ser Asp

Gly Asn Gly Pro Thr Gly Asn Gly Asp His Pro Asn Phe Ser Leu Ser 20 . 25

Glu Gly Arg Ala Phe Gly Ser Leu Ala Ala Gln Pro Ile Thr Ser Cys 35 - 40

Leu Ser Val Pro Ala Pro Pro Phe Ser Leu Gly Lys Leu Gln Asp Gly

Leu Leu His Ile Thr Thr Cys Ser Phe Val Ala Pro Trp Asn Ser Leu

Ser Leu Ala Gln Arg Arg Gly Phe Thr Lys Thr Tyr Thr Val Gly Cys

Glu Glu Cys Thr Val Phe Pro Cys Leu Ser Ile Pro Cys Lys Leu Gln 105

Ser Gly Thr His Cys Leu Trp Thr Asp Gln Leu Leu Gln Gly Ser Glu 120

Lys Gly Phe Gln Ser Arg His Leu Ala Cys Leu Pro Arg Glu Pro Gly

Leu Cys Thr Trp Gln Ser Leu Arg Ser Gln Ile Ala 145 150

<210> 238 <211> 226

<212> PRT

<213> Homo sapien

<400> 238

Met Ala Ala Ala Ala Ala Ala Gly Ala Ala Gly Ser Ala Ala Pro 10

Ala Ala Ala Gly Ala Pro Gly Ser Gly Gly Ala Pro Ser Gly Ser

Gln Gly Val Leu Ile Gly Asp Arg Leu Tyr Ser Gly Val Leu Ile Thr 40

Leu Glu Asn Cys Leu Leu Pro Asp Asp Lys Leu Arg Phe Thr Pro Ser 50

Met Ser Ser Gly Leu Asp Thr Asp Thr Glu Thr Asp Leu Arg Val Val 70

Gly Cys Glu Leu Ile Gln Ala Ala Gly Ile Leu Leu Arg Leu Pro Gln 85 90 95

Val Ala Met Ala Thr Gly Gln Val Leu Phe Gln Arg Phe Phe Tyr Thr 100 105

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Lys Ser Phe Val Lys His Ser Met Glu His Val Ser Met Ala Cys Val 115 120

His Leu Ala Ser Lys Ile Glu Glu Ala Pro Arg Arg Ile Arg Asp Val

Ile Asn Val Phe His Arg Leu Arg Gln Leu Arg Asp Lys Lys Pro

Val Pro Leu Leu Asp Gln Asp Tyr Val Asn Leu Lys Asn Gln Ile 165 170

Ile Lys Ala Glu Arg Arg Val Leu Lys Glu Leu Gly Phe Cys Val His 180 185 190

Val Lys His Pro His Lys Ile Ile Val Met Tyr Leu Gln Val Leu Glu 195

Cys Glu Arg Asn Gln His Leu Val Gln Thr Ser Trp Val Ala Ser Glu 210 215

Gly Lys 225

<210> 239

<211> 253 <212> PRT

<213> Homo sapien

<400> 239

Asp Ser Gln Asp Cys Leu Ala Leu Ser Pro Ser Asn Arg Leu Leu Arg

Gly Val Val Arg Leu Ser Arg Phe Ser Leu Asp Asn Ala Gly Gly Arg 25

Pro Gly Phe Pro Gly Gly Ala Leu Gln Phe Phe Leu Cys Leu Ala Ser

Arg Asn Tyr Met Asn Asp Ser Leu Arg Thr Asp Val Phe Val Arg Phe

Gln Pro Glu Ser Ile Ala Cys Ala Cys Ile Tyr Leu Ala Ala Arg Thr

Leu Glu Ile Pro Leu Pro Asn Arg Pro His Trp Phe Leu Leu Phe Gly 90

Ala Thr Glu Glu Glu Ile Gln Glu Ile Cys Leu Lys Ile Leu Gln Leu 100 105 110

Tyr Ala Arg Lys Lys Val Asp Leu Thr His Leu Glu Gly Glu Val Glu 115 120 125

Lys Arg Lys His Ala Ile Glu Glu Ala Lys Ala Gln Ala Arg Gly Leu 130 135 140

Leu Pro Gly Gly Thr Gln Val Leu Asp Gly Thr Ser Gly Phe Ser Pro 145 150 155 160

Ala Pro Lys Leu Val Glu Ser Pro Lys Glu Gly Lys Gly Ser Lys Pro 165 170 175

Ser Pro Leu Ser Val Lys Asn Thr Lys Arg Arg Leu Glu Gly Ala Lys 180 185 190

Lys Ala Lys Ala Asp Ser Pro Val Asn Gly Leu Pro Lys Gly Arg Glu 195 200 205

Ser Arg Ser Arg Ser Arg Ser Arg Glu Gln Ser Tyr Ser Arg Ser Pro 210 215 220

Ser Arg Ser Ala Ser Pro Lys Arg Arg Lys Ser Asp Ser Gly Ser Thr 225 230 235 240

Ser Gly Gly Ser Lys Ser Gln Arg Ser Leu Gln Arg Leu 245 250

<210> 240

<211> 346

<212> PRT

<213> Homo sapien

<400> 240

Asp Ser Gln Asp Cys Leu Ala Leu Ser Pro Ser Asn Arg Leu Leu Arg 1 5 10 15

Gly Val Val Arg Leu Ser Arg Phe Ser Leu Asp Asn Ala Gly Gly Arg

Pro Gly Phe Pro Gly Gly Ala Leu Gln Phe Phe Leu Cys Leu Ala Ser 35 40

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Arg Asn Tyr Met Asn Asp Ser Leu Arg Thr Asp Val Phe Val Arg Phe 50 55 60

Gln Pro Glu Ser Ile Ala Cys Ala Cys Ile Tyr Leu Ala Ala Arg Thr 65 70 75 80

Leu Glu Ile Pro Leu Pro Asn Arg Pro His Trp Phe Leu Leu Phe Gly 85 90 95

Ala Thr Glu Glu Glu Ile Gln Glu Ile Cys Leu Lys Ile Leu Gln Leu 100 105 110

Tyr Ala Arg Lys Lys Val Asp Leu Thr His Leu Glu Gly Glu Val Glu 115 120 125

Lys Arg Lys His Ala Ile Glu Glu Ala Lys Ala Gln Ala Arg Gly Leu 130 135 140

Leu Pro Gly Gly Thr Gln Val Leu Asp Gly Thr Ser Gly Phe Ser Pro 145 155 . 160

Ala Pro Lys Leu Val Glu Ser Pro Lys Glu Gly Lys Gly Ser Lys Pro 165 170 175

Ser Pro Leu Ser Val Lys Asn Thr Lys Arg Arg Leu Glu Gly Ala Lys 180 185 190

Lys Ala Lys Ala Asp Ser Pro Val Asn Gly Leu Pro Lys Gly Arg Glu
195 200 205

Ser Arg Ser Arg Ser Arg Ser Arg Glu Gln Ser Tyr Ser Arg Ser Pro 210 215 220

Ser Arg Ser Ala Ser Pro Lys Arg Arg Lys Ser Asp Ser Gly Ser Thr 225 230 235 240

Ser Gly Gly Ser Lys Ser Gln Ser Arg Ser Arg Ser Arg Ser Asp Ser 245 250 255

Pro Pro Arg Glm Ala Pro Arg Ser Ala Pro Tyr Lys Gly Ser Glu Ile 260 265 270

Arg Gly Ser Arg Lys Ser Lys Asp Cys Lys Tyr Pro Gln Lys Pro His 275 280 285

Lys Ser Arg Ser Arg Ser Ser Ser Arg Ser Arg Ser Arg Ser Arg Glu

290 295 300

Arg Ala Asp Asn Pro Gly Lys Tyr Lys Lys Lys Ser His Tyr Tyr Arg 305 310 315 320

Asp Gln Arg Arg Glu Arg Ser Arg Ser Tyr Glu Arg Thr Gly Arg Arg 325 330 335

Tyr Glu Arg Asp His Pro Val Ala Ala Leu 340 345

<210> 241

<211> 91

<212> PRT

<213> Homo sapien

<400> 241

Pro Thr Thr Thr Lys Phe Ala Ala Ala Ser Thr Phe Leu Asn Trp Cys
1 5 10 15

Cys Leu Gly Phe Ile Ala Phe Ala Tyr Ser Val Lys Ser Arg Asp Arg 20 25 30

Lys Met Val Gly Asp Val Thr Gly Ala Gln Ala Tyr Ala Ser Thr Ala 35 40 45

Lys Cys Leu Asn Ile Trp Ala Leu Ile Leu Gly Ile Phe Met Thr Ile 50 55

Gly Phe Ile Leu Leu Val Phe Gly Ser Val Thr Val Tyr His Ile 65 70 75 80

Met Leu Gln Ile Ile Gln Glu Lys Arg Gly Tyr 85 90

<210> 242

<211> 92

<212> PRT

<213> Homo sapien

<400> 242

Gly Gln Glu Asp Gly Trp Arg Arg Asp Arg Gly Pro Gly Leu Cys Leu 1 5 10 15

His Arg Gln Val Pro Glu His Leu Gly Pro Asp Phe Gly His Leu His 20 25 30

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Asp Arg Ile His Pro Val Thr Gly Ile Arg Leu Cys Asp Ser Leu Pro 40

Tyr Tyr Val Thr Asp Asn Thr Gly Lys Thr Gly Leu Leu Val Ala Ala 50

His Ser Leu Gln Pro Leu His Ser Thr Val Gln Cys Trp Pro Cys Thr 65 70

Leu Gly Leu Leu Pro Leu Pro Pro Trp Ser Cys Pro

<210> 243 <211> 137 <212> PRT <213> Homo sapien

<400> 243

Met Leu Glu Arg Arg Ser Val Met Asp Arg Pro Pro Ala Glu Val Arg 10

Glu Thr Lys Ile Lys Gly Lys Ser Gly Arg Phe Phe Thr Val Lys Leu 25

Pro Val Ala Leu Asp Pro Gly Ala Lys Ile Ser Val Ile Val Glu Thr

Val Tyr Thr His Val Leu His Pro Tyr Pro Thr Gln Ile Thr Gln Ser 55

Glu Lys Gln Phe Val Val Phe Glu Gly Asn His Tyr Phe Tyr Ser Pro

Tyr Pro Thr Lys Thr Gln Thr Met Arg Val Lys Leu Ala Ser Arg Asn

Val Glu Ser Tyr Thr Lys Leu Gly Asn Pro Thr Arg Ser Glu Asp Leu 100 105

Leu Asp Tyr Gly Pro Phe Arg Asp Val Pro Ala Tyr Ser Gln Asp Thr 115 · 120

Phe Lys Val Pro Arg Pro Arg Pro Arg 130

<210> 244 <211> 148

••

<212> PRT

<213> Homo sapien

<220>

<221> MISC FEATURE

<222> (21)..(22)

<223> X=any amino acid

<220>

<221> MISC\_FEATURE '

<222> (24)..(24) <223> X=any amino acid

<400> 244

Arg Leu Ile Tyr Arg Ala Ile Gly His Leu Ile Met Leu Glu Arg Arg

Ser Val Met Asp Xaa Xaa Pro Xaa Glu Val Arg Glu Thr Lys Ile Lys 20

Gly Lys Ser Gly Arg Phe Phe Thr Val Lys Leu Pro Val Ala Leu Aso 40

Pro Gly Ala Lys Ile Ser Val Ile Val Glu Thr Val Tyr Thr His Val 55

Leu His Pro Tyr Pro Thr Gln Ile Thr Gln Ser Glu Lys Gln Phe Val

Val Phe Glu Gly Asn His Tyr Phe Tyr Ser Pro Tyr Pro Thr Lys Thr 85

Gln Thr Met Arg Val Lys Leu Ala Ser Arg Asn Val Glu Ser Tyr Thr

Lys Leu Gly Asn Pro Thr Arg Ser Glu Asp Leu Leu Asp Tyr Gly Pro

Phe Arg Asp Val Pro Ala Tyr Ser Gln Asp Thr Phe Lys Val Pro Arg 130 135

Pro Arg Pro Arg 145

<210> 245

<211> 479

<212> PRT

<213> Homo sapien

. <400> 245

Met Glu Ala Pro Ala Ala Gly Leu Phe Leu Leu Leu Leu Gly Thr 1 5 10 15

Trp Ala Pro Ala Pro Gly Ser Ala Ser Ser Glu Ala Pro Pro Leu Ile 20 25 30

Asn Glu Asp Val Lys Arg Thr Val Asp Leu Ser Ser His Leu Ala Lys 35 40 45

Val Thr Ala Glu Val Val Leu Ala His Leu Gly Gly Ser Thr Ser 50 55

Arg Ala Thr Ser Phe Leu Leu Ala Leu Glu Pro Glu Leu Glu Ala Arg 65 70 75 80

Leu Ala His Leu Gly Val Glm Val Lys Gly Glu Asp Glu Glu Glu Asn 85 90 95

Asn Leu Glu Val Arg Glu Thr Lys Ile Lys Gly Lys Ser Gly Arg Phe 100 105 110

Phe Thr Val Lys Leu Pro Val Ala Leu Asp Pro Gly Ala Lys Ile Ser

Val Ile Val Glu Thr Val Tyr Thr His Val Leu His Pro Tyr Pro Thr 130 135 140

Gln Ile Thr Gln Ser Glu Lys Gln Phe Val Val Phe Glu Gly Asn His 145 150 150 155

Tyr Phe Tyr Ser Pro Tyr Pro Thr Lys Thr Gln Thr Met Arg Val Lys 165 170 175

Leu Ala Ser Arg Asn Val Glu Ser Tyr Thr Lys Leu Gly Asn Pro Thr 180 185 190

Arg Ser Glu Asp Leu Leu Asp Tyr Gly Pro Phe Arg Asp Val Pro Ala 195 200 205

Tyr Ser Gln Asp Thr Phe Lys Val His Tyr Glu Asn Asn Ser Pro Phe 210 220

Leu Thr Ile Thr Ser Met Thr Arg Val Ile Glu Val Ser His Trp Gly 225 230 235 240

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Asn Ile Ala Val Glu Glu Asn Val Asp Leu Lys His Thr Gly Ala Val

Leu Lys Gly Pro Phe Ser Arg Tyr Asp Tyr Gln Arg Gln Pro Asp Ser 265

Gly Ile Ser Ser Ile Arg Ser Phe Lys Thr Ile Leu Pro Ala Ala Ala 280

Gln Asp Val Tyr Tyr Arg Asp Glu Ile Gly Asn Val Ser Thr Ser His 295

Leu Leu Ile Leu Asp Asp Ser Val Glu Met Glu Ile Arg Pro Arg Phe 315 305 310

Pro Leu Phe Gly Gly Trp Lys Thr His Tyr Ile Val Gly Tyr Asn Leu

Pro Ser Tyr Glu Tyr Leu Tyr Asn Leu Gly Asp Gln Tyr Ala Leu Lys 345

Met Arg Phe Val Asp His Val Phe Asp Glu Gln Val Ile Asp Ser Leu 360 355

Thr Val Lys Ile Ile Leu Pro Glu Gly Ala Lys Asn Ile Glu Ile Asp

Ser Pro Tyr Glu Ile Ser Arg Ala Pro Asp Glu Leu His Tyr Thr Tyr 395

Leu Asp Thr Phe Gly Arg Pro Val Ile Val Ala Tyr Lys Lys Asn Leu 410 405

Val Glu Gln His Ile Gln Asp Ile Val Leu Asp Ala Gln Val Lys Glu 425 420

Leu Val Leu Lys Ser Ala Val Glu Ala Glu Arg Leu Val Ala Gly Lys 440

Leu Lys Lys Asp Thr Tyr Ile Glu Asn Glu Lys Leu Ile Ser Gly Lys

Arg Gln Glu Leu Val Thr Lys Ile Asp His Ile Leu Asp Ala Leu 470

<210> 246

<211> 361 <212> PRT

<213> Homo sapien

<400> 246

Met Glu Ala Pro Ala Ala Gly Leu Phe Leu Leu Leu Leu Leu Gly Thr 1 5 10 15

Trp Ala Pro Ala Pro Gly Ser Ala Ser Ser Glu Ala Pro Pro Leu Ile 20 25 30

Asn Glu Asp Val Lys Arg Thr Val Asp Leu Ser Ser His Leu Ala Lys 35 40 45

Val Thr Ala Glu Val Val Leu Ala His Leu Gly Gly Ser Thr Ser 50 55 60

Arg Ala Thr Ser Phe Leu Leu Ala Leu Glu Pro Glu Leu Glu Ala Arg 65 70 75 80

Leu Ala His Leu Gly Val Gln Val Lys Gly Glu Asp Glu Glu Glu Asn 85 90 95

Asn Leu Glu Val Arg Glu Thr Lys Ile Lys Gly Lys Ser Gly Arg Phe 100 105 110

Phe Thr Val Lys Leu Pro Val Ala Leu Asp Pro Gly Ala Lys Ile Ser 115 120 125

Val Ile Val Glu Thr Val Tyr Thr His Val Leu His Pro Tyr Pro Thr 130 135 140

Gln Ile Thr Gln Ser Glu Lys Gln Phe Val Val Phe Glu Gly Asn His 145 150 155 160

Tyr Phe Tyr Ser Pro Tyr Pro Thr Lys Thr Gln Thr Met Arg Val Lys
165 170 175

Leu Ala Ser Arg Asn Val Glu Ser Tyr Thr Lys Leu Gly Asn Pro Thr 180 185 190

Arg Ser Glu Asp Leu Leu Asp Tyr Gly Pro Phe Arg Asp Val Pro Ala 195 200 205

Tyr Ser Gln Asp Thr Phe Lys Val His Tyr Glu Asn Asn Ser Pro Phe

220 210 215

Leu Thr Ile Thr Ser Met Thr Arg Val Ile Glu Val Ser His Trp Gly 235 230

Asn Ile Ala Val Glu Glu Asn Val Asp Leu Lys His Thr Gly Ala Val 250

Leu Lys Gly Pro Phe Ser Arg Tyr Asp Tyr Gln Arg Gln Pro Asp Ser 265 260

Gly Ile Ser Ser Ile Arg Ser Phe Lys Thr Ile Leu Pro Ala Ala Ala 280 .

Gln Asp Val Tyr Tyr Arg Asp Glu Ile Gly Asn Val Ser Thr Ser His

Leu Leu Ile Leu Asp Asp Ser Val Glu Met Glu Ile Arg Pro Arg Phe

Pro Leu Phe Gly Gly Trp Lys Thr His Tyr Ile Val Gly Tyr Asn Leu 330

Pro Ser Tyr Glu Tyr Leu Tyr Asn Leu Gly Gln Ser Ser Ile Val Arg

Glu Lys Leu Thr Phe Ser Leu Ile Ser

<210> 247 <211> 420 <212> PRT <213> Homo sapien

<400> 247

Met Glu Ala Pro Ala Ala Gly Leu Phe Leu Leu Leu Leu Gly Thr 5

Trp Ala Pro Ala Pro Gly Ser Ala Ser Ser Glu Ala Pro Pro Leu Ile 20 25 30

Asn Glu Asp Val Lys Arg Thr Val Asp Leu Ser Ser His Leu Ala Lys 40 . 45 35

Val Thr Ala Glu Val Val Leu Ala His Leu Gly Gly Ser Thr Ser 50 55

Arg Ala Thr Ser Phe Leu Leu Ala Leu Glu Pro Glu Leu Glu Ala Arg 75 70 Leu Ala His Leu Gly Val Gln Val Lys Gly Glu Asp Glu Glu Glu Asn 85 Asn Leu Glu Val Arg Glu Thr Lys Ile Lys Gly Lys Ser Gly Arg Phe Phe Thr Val Lys Leu Pro Val Ala Leu Asp Pro Gly Ala Lys Ile Ser 120 Val Ile Val Glu Thr Val Tyr Thr His Val Leu His Pro Tyr Pro Thr 135 Gln Ile Thr Gln Ser Glu Lys Gln Phe Val Val Phe Glu Gly Asn His 150 Tyr Phe Tyr Ser Pro Tyr Pro Thr Lys Thr Gln Thr Met Arg Val Lys 170 Leu Ala Ser Arg Asn Val Glu Ser Tyr Thr Lys Leu Gly Asn Pro Thr 180 185 Arg Ser Glu Asp Leu Leu Asp Tyr Gly Pro Phe Arg Asp Val Pro Ala 195 200 205 Tyr Ser Gln Asp Thr Phe Lys Val His Tyr Glu Asn Asn Ser Pro Phe 215 Leu Thr Ile Thr Ser Met Thr Arg Val Ile Glu Val Ser His Trp Gly 235 Asn Ile Ala Val Glu Glu Asn Val Asp Leu Lys His Thr Gly Ala Val 250 Leu Lys Gly Pro Phe Ser Arg Tyr Asp Tyr Gln Arg Gln Pro Asp Ser 265 Gly Ile Ser Ser Ile Arg Ser Phe Lys Thr Ile Leu Pro Ala Ala Ala

Gln Asp Val Tyr Tyr Arg Asp Glu Ile Gly Asn Val Ser Thr Ser His

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Leu Leu Ile Leu Asp Asp Ser Val Glu Met Glu Ile Arg Pro Arg Phe 305 310 315 320

Pro Leu Phe Gly Gly Trp Lys Thr His Tyr Ile Val Gly Tyr Asn Leu 325 330 335

Pro Ser Tyr Glu Tyr Leu Tyr Asn Leu Gly Asp Gln Tyr Ala Leu Lys 340 345 350

Met Arg Phe Val Asp His Val Phe Asp Glu Gln Val Ile Asp Ser Leu 355 360 365

Thr Val Lys Ile Ile Leu Pro Glu Gly Ala Lys Thr Thr Ser Trp Met 370 380

Pro Cys Ser Pro Cys Pro His Pro Pro Gly Gly Pro Gly Cys Leu His 385 390 395 400

Phe Ala Val Ala Gly Arg Leu Gly Gly Ser Gly Arg Leu Cys Met Glu 405 415

Ala Ser Glu Ser 420

<210> 248

<211> 128

<212> PRT

<213> Homo sapien

<400> 248

Gly Cys Ala Arg Glu Pro Glu Ser Arg Leu Pro Lys Leu Gly Ser Trp 1 5 10 15

Glu Asn Leu Gly Pro Gly Leu Thr Glu Lys Arg Arg Gly Lys Glu Ala 20 25 30

Gly Gln Glu Gly Ala Trp Arg Thr Pro Ala Gly Gly Arg Gly Ala 35 40 45

Ala Gly Leu Ser Val Thr Pro Leu Ser Pro Pro Arg Pro Ala Pro Pro 50 55 60

Ala Gly Glu Gly Pro Arg Cys Pro Pro Gly Arg Pro Ala Pro Ala Arg 65 70 75 80

Arg Arg Lys Gly Trp Arg Val Glu Arg Gly Gly Arg Gly Ser Ala 85 90 95

Trp Asp Ala Pro Gly His Arg Ala Arg Ser Leu Arg Pro Gly Ala Gly 100 105 110

Gln Val Arg Gly Gln Asp Val Gly Arg Thr Trp Cys Met Ala Ala Ser 115 120 125

<210> 249

<211> 315

<212> PRT

<213> Homo sapien

<400> 249

Met Ser Ala Ala Gly Ala Gly Ala Gly Val Glu Ala Gly Phe Ser Ser 1 5 10 15

Glu Glu Leu Leu Ser Leu Arg Phe Pro Leu His Arg Ala Cys Arg Asp 20 25 30

Gly Asp Leu Ala Thr Leu Cys Ser Leu Leu Gln Gln Thr Pro His Ala 35 40 45

His Leu Ala Ser Glu Asp Ser Phe Tyr Gly Trp Thr Pro Val His Trp 50 60

Ala Ala His Phe Gly Lys Leu Glu Cys Leu Val Gln Leu Val Arg Ala 65 70 75 80

Gly Ala Thr Leu Asn Val Ser Thr Thr Arg Tyr Ala Gln Thr Pro Ala 85 90 95

His Ile Ala Ala Phe Gly Gly His Pro Gln Cys Leu Val Trp Leu Ile 100 105 110

Gln Ala Gly Ala Asn Ile Asn Lys Pro Asp Cys Glu Gly Glu Thr Pro 115 120 125

Ile His Lys Ala Ala Arg Ser Gly Ser Leu Glu Cys Ile Ser Ala Leu 130 135 140

Val Ala Asn Gly Ala His Val Asp Leu Arg Asn Ala Ser Gly Leu Thr 145 150 155 160

Ala Ala Asp Ile Ala Gln Thr Gln Gly Phe Gln Glu Cys Ala Gln Phe

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Leu Leu Asn Leu Gln Asn Cys His Leu Asn His Phe Tyr Asn Asn Gly 180 185 190

Ile Leu Asn Gly Gly His Gln Asn Val Phe Pro Asn His Ile Ser Val 195 200 205

Gly Thr Asn Arg Lys Arg Cys Leu Glu Asp Ser Glu Asp Phe Gly Val 210 215 220

Lys Lys Ala Arg Thr Glu Ala Gln Ser Leu Asp Ser Ala Val Pro Leu 225 230 235 240

Thr Asn Gly Asp Thr Glu Asp Asp Ala Asp Lys Met His Val Asp Arg 245 250 255

Glu Phe Ala Val Val Thr Gly Gly Ser Gly Gln Phe Pro Val Ser Cys 260 265 270

Asn Asn Pro Met Val Glu Asp Thr Lys Gln Gln Glu Ser Gly Ser 275 280 285

Val Gly Pro Lys Glu Ile Glu Ile Tyr Thr Val Ser Ala Met Gln Thr 290 295 300

Pro Cys Arg Cys Arg Asn Gln Tyr Glu Lys Gln 305 310 315

<210> 250

<211> 142

<212> PRT

<213> Homo sapien

<400> 250

Met Gly Asn Phe Lys Ser Ile Ser Thr Ser Thr Lys Met Val Asn Gly

1 10 15

Arg Lys Ile Thr Thr Lys Arg Ile Val Glu Asn Gly Gln Glu Arg Val 20 25 30

Glu Val Glu Glu Asp Gly Gln Leu Lys Ser Leu Thr Ile Asn Gly Val 35 40 45

Ala Asp Asp Asp Ala Leu Ala Glu Glu Arg Met Arg Arg Gly Gln Asn 50 55 60

Ala Leu Pro Ala Gln Pro Ala Gly Leu Arg Pro Pro Lys Pro Pro Arg 65 70 75 80

Pro Ala Ser Leu Leu Arg His Ala Pro His Cys Leu Ser Glu Glu Glu 85 90 95

Gly Glu Gln Asp Arg Pro Arg Ala Pro Gly Pro Trp Asp Pro Leu Ala 100 105 110

Ser Ala Ala Gly Leu Lys Glu Gly Gly Lys Arg Lys Lys Gln Lys Gln 115 120 125

Arg Glu Glu Ser Lys Lys Lys Ser Thr Lys Gly Asn His 130 135 140

<210> 251

<211> 72

<212> PRT

<213> Homo sapien

<400> 251

Met Gly Leu Ser His Ala Gly Trp His Arg Ala Gly Lys His Glu Ala 1 5 10 15

Ser Pro His Gln Gly Phe Ala Cys Arg Lys Ala Ala Leu Trp Pro Ala 20 25 30

Gly Glu Ala Glu Glu Thr Pro Val Asp Thr Leu Pro Thr Gly Leu Lys 35 40 45

Glu Gly Gly Lys Arg Lys Lys Gln Lys Gln Arg Glu Glu Ser Lys Lys 50 55

Lys Lys Ser Thr Lys Gly Asn His 65 70

<210> 252

<211> 122

<212> PRT

<213> Homo sapien

<400> 252

Thr Gly Leu Glù Ala Arg Gly Ala Pro Pro Asp Ala Gly Ala Pro Pro 1 5 10 15

Cys Ser Ala Cys Gly Arg Ala His Ala Leu Gly Ser Ser Val Gly Gln 20 25 30

Asp Cys Leu Glu Ala Thr Leu Ala Arg Gln Asp Tyr Ala Ile Thr Asp

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40 45 35

Gln Ser Glu Gln Gly Gln Glu Thr Gly Leu Thr Ala Arg Val Ala Gly

Thr Asp Val Trp Asp Leu Ala Ala Thr Leu Cys Phe Ser Pro Ala Leu 70

Asn Leu Leu His Phe Pro Leu Val Leu Pro Asp Pro Leu His Ser Phe 90 85

Arg Leu Leu Asn His Ser Ala Cys Cys Trp Asn Ile Ser Gly Phe Arg 100 105

Ser Thr Gly Gly Arg Arg Trp Leu Thr Glu

. <210> 253

<211> 42

<212> PRT

<213> Homo sapien

<400> 253

Met Ala Lys Lys Ala Gly Leu Cys Leu Gly Gly Ser Arg Gln Gly Gly 1 5 10 15

Cys Gln Ser Gly Met Val Thr Gly Asn Glu Pro Arg Asp Leu Ala Leu 20 25 30

Ser His Pro Leu Ser Phe Val Gly Gly Leu

<210> 254 <211> 260 <212> PRT <213> Homo sapien

<400> 254

Val Phe Cys Ser Phe Phe Ala Glu Lys Glu Gln Gln Glu Ala Ile Glu

His Ile Asp Glu Val Gln Asn Glu Ile Asp Arg Leu Asn Glu Gln Ala 20 25

Ser Glu Glu Ile Leu Lys Val Glu Gln Lys Tyr Asn Lys Leu Arg Gln 40

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Pro Phe Phe Gln Lys Arg Ser Glu Leu Ile Ala Lys Ile Pro Asn Phe 55

Trp Val Thr Thr Phe Val Asn His Pro Gln Val Ser Ala Leu Leu Gly 75 70

Glu Glu Asp Glu Glu Ala Leu His Tyr Leu Thr Arg Val Glu Val Thr 85

Glu Phe Glu Asp Ile Lys Ser Gly Tyr Arg Ile Asp Phe Tyr Phe Asp 100

Glu Asn Pro Tyr Phe Glu Asn Lys Val Leu Ser Lys Glu Phe His Leu 120

Asn Glu Ser Gly Asp Pro Ser Ser Lys Ser Thr Glu Ile Lys Trp Lys 135

Ser Gly Lys Asp Leu Thr Lys Arg Ser Ser Gln Thr Gln Asn Lys Ala 150

Ser Arg Lys Arg Gln His Glu Glu Pro Glu Ser Phe Phe Thr Trp Phe 170

Thr Asp His Ser Asp Ala Gly Ala Asp Glu Leu Gly Glu Val Ile Lys 180 185

Asp Asp Ile Trp Pro Asn Pro Leu Gln Tyr Tyr Leu Val Pro Asp Met 200 195

Asp Asp Glu Glu Gly Glu Glu Glu Asp Asp Asp Asp Glu Glu

Glu Glu Gly Leu Glu Asp Ile Asp Glu Glu Gly Asp Glu Asp Glu Gly 235

Glu Glu Asp Glu Asp Asp Asp Glu Gly Glu Glu Glu Glu Asp Glu 250 245

Gly Glu Asp Asp 260

<210> 255 <211> 285 <212> PRT <213> Homo sapien

| <400> | 255 |
|-------|-----|
|-------|-----|

Ser Leu Gln Asp Lys Arg Ala Pro Ile Pro Glu His Thr Pro Phe Ser 1 5 10 15

Ser Ser Pro Phe Cys Ala Ser Leu Leu Ser Asp Leu Ile Val Ala Pro  $20 \hspace{1cm} 25 \hspace{1cm} 30$ 

Lys Lys Glu Gln Gln Glu Ala Ile Glu His Ile Asp Glu Val Gln Asn  $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$ 

Glu Gln Lys Tyr Asn Lys Leu Arg Gln Pro Phe Phe Gln Lys Arg Ser 65 70 75 80

Glu Leu Ile Ala Lys Ile Pro Asn Phe Trp Val Thr Thr Phe Val Asn 85 90 95

His Pro Gln Val Ser Ala Leu Leu Gly Glu Glu Asp Glu Glu Ala Leu 100 105 110

His Tyr Leu Thr Arg Val Glu Val Thr Glu Phe Glu Asp Ile Lys Ser 115 120 125

Gly Tyr Arg Ile Asp Phe Tyr Phe Asp Glu Asn Pro Tyr Phe Glu Asn 130 135

Lys Val Leu Ser Lys Glu Phe His Leu Asn Glu Ser Gly Asp Pro Ser 145 150 155 160

Ser Lys Ser Thr Glu Ile Lys Trp Lys Ser Gly Lys Asp Leu Thr Lys 165 170 175

Arg Ser Ser Gln Thr Gln Asn Lys Ala Ser Arg Lys Arg Gln His Glu 180 185 190

Glu Pro Glu Ser Phe Phe Thr Trp Phe Thr Asp His Ser Asp Ala Gly
195 200 205

Leu Gln Tyr Tyr Leu Val Pro Asp Met Asp Asp Glu Glu Gly Glu Gly 225 230 235 240

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Glu Glu Asp Asp Asp Asp Glu Glu Glu Glu Gly Leu Glu Asp Ile
245 250 255

Asp Glu Glu Gly Asp Glu Asp Glu Gly Glu Asp Glu Asp Asp Asp 260 265 270

Glu Gly Glu Gly Glu Glu Gly Glu Gly Glu Asp Asp 275 280 285

<210> 256

<211> 600

<212> PRT

<213> Homo sapien

<400> 256

Met Ala Thr Pro Leu Pro Gly Arg Ala Gly Gly Pro Ala Thr Pro Leu 1 5 10 15

Ser Pro Thr Arg Leu Ser Arg Leu Gln Glu Lys Glu Glu Leu Arg Glu 20 25 30

Leu Asn Asp Arg Leu Ala His Tyr Ile Asp Arg Val Arg Ala Leu Glu 35 40 45

Leu Glu Asn Asp Arg Leu Leu Leu Lys Ile Ser Glu Lys Glu Glu Val 50 55 60

Thr Thr Arg Glu Val Ser Gly Ile Lys Ala Leu Tyr Glu Ser Glu Leu 65 70 75 80

Ala Asp Ala Arg Val Leu Asp Glu Thr Ala Arg Glu Arg Ala Arg 85 90 95

Leu Gln Ile Glu Ile Gly Lys Leu Arg Ala Glu Leu Asp Glu Val Asn 100 105 110

Lys Ser Ala Lys Lys Arg Glu Gly Glu Leu Thr Val Ala Gln Gly Arg 115 120 125

Val Lys Asp Leu Glu Ser Leu Phe His Arg Ser Glu Val Glu Leu Ala 130 135 140

Ala Ala Leu Ser Asp Lys Arg Gly Leu Glu Ser Asp Val Ala Glu Leu 145 150 155 160

Arg Ala Gln Leu Ala Lys Ala Glu Asp Gly His Ala Val Ala Lys Lys

|            |             |            | 328          |              |            |              |              |            |            |            |              |            |            |                    |            |
|------------|-------------|------------|--------------|--------------|------------|--------------|--------------|------------|------------|------------|--------------|------------|------------|--------------------|------------|
|            |             |            |              | 165          |            |              |              |            | 170        |            |              |            |            | 175                |            |
| Gln        | Leu         | Glu        | Lys<br>180   | Glu          | Thr        | Leu          | Met          | Arg<br>185 | Val        | Asp        | Leu          | Glu        | Asn<br>190 | Arg                | Cys        |
| Gln        | Ser         | Leu<br>195 | Gln          | Glu          | Glu        | Leu          | Asp<br>200   | Phe        | Arg        | Lys        | Ser          | Val<br>205 | Phe        | Glu                | Glu        |
| Glu        | Val<br>210  | Arg        | Glu          | Thr          | Arg        | Arg<br>215   | Arg          | His        | Glu        | Arg        | Arg<br>220   | Leu        | Val        | Glu                | Val        |
| Asp<br>225 | Ser         | Ser        | Arg          | Gln          | Gln<br>230 | Glu          | Tyr          | Asp        | Phe        | Lys<br>235 | Met          | Ala        | Gln        | Ala                | Leu<br>240 |
| Glu        | Glu         | Leu        | Arg          | Ser<br>245   | Gln        | His          | Asp          | Glu        | Gln<br>250 | Val        | Arg          | Leu        | Tyr        | <u>Б</u> уs<br>255 | Leu        |
| Glu        | Leu         | Glu        | Gln<br>260   |              | Tyr        | Gln          | Ala          | Lys<br>265 | Leu        | Asp        | Ser          | Ala        | Lys<br>270 | Leu                | Ser        |
| Ser        | Asp         | Gln<br>275 |              | qeA .        | Lys        | Ala          | Ala<br>280   | Ser        | Ala        | Ala        | Arg          | Glu<br>285 | Glu        | Leu                | Lys        |
| Glu        | Ala<br>290  |            | , Met        | Arg          | Leu        | Glu<br>295   |              | Leu        | Ser        | Tyr        | Gln<br>300   | Leu        | Ser        | Gly                | Leu        |
| Gln<br>305 |             | Glr        | n Ala        | Ser          | Ala<br>310 |              | . Glu        | qsA        | Arg        | Ile<br>315 | Arg          | Glu        | Leu        | . Glu              | Glu<br>320 |
| Ala        | . Met       | : Ala      | a Gly        | 7 Glu<br>325 |            | Asp          | Lys          | Phe        | Arg<br>330 |            | Met          | Leu        | . Asp      | Ala<br>335         | Lys        |
| Glu        | ı Glr       | ı Glı      | 1 Met<br>340 |              | Glu        | ı Met        | : Arg        | Asp<br>345 |            | . Met      | . Gln        | Gln        | 350        | ı Leu<br>)         | Ala        |
| Glı        | 1 Ту        | Gl:<br>35  |              | ı Let        | ı Lev      | ı Asp        | y Val<br>360 |            | s Lev      | ı Ala      | . Leu        | Asg<br>365 | Met        | : Glu              | ı Ile      |
| Ası        | n Ala<br>37 |            | r Ar         | g Ly         | s Lev      | ı Let<br>379 |              | ı Gly      | y Glu      | ı Glu      | 1 Glu<br>380 | 1 Arg      | g Le       | ı Lys              | s Leu      |
| <br>Se:    | r Pr        | o Se       | r Pr         | o Se         | r Se       | r Ár         | g Vai        | l Th       | r Val      | l Sei      | r Arg        | Älä        | a Th:      | r Sei              | r Ser      |

Ser Ser Gly Ser Leu Ser Ala Thr Gly Arg Leu Gly Arg Ser Lys Arg

Lys Arg Leu Glu Val Glu Glu Pro Leu Gly Ser Gly Pro Ser Val Leu 420 425 430

Gly Thr Gly Gly Ser Gly Gly Phe His Leu Ala Gln Gln Ala 435 440 445

Ser Ala Ser Gly Ser Val Ser Ile Glu Glu Ile Asp Leu Glu Gly Lys 450 455 460

Phe Val Gln Leu Lys Asn Asn Ser Asp Lys Asp Gln Ser Leu Gly Asn 465 470 475 480

Trp Arg Ile Lys Arg Gln Val Leu Glu Glu Glu Glu Ile Ala Tyr Lys 485 490 495

Phe Thr Pro Lys Tyr Ile Leu Arg Ala Gly Gln Met Val Thr Val Trp
500 505 510

Ala Ala Gly Ala Gly Val Ala His Ser Pro Pro Ser Thr Leu Val Trp  $515 \hspace{1cm} 520 \hspace{1cm} 525 \hspace{1cm}$ 

Lys Gly Gln Ser Ser Trp Gly Thr Gly Glu Ser Phe Arg Thr Val Leu 530 540

Val Asn Ala Asp Gly Glu Glu Val Ala Met Arg Thr Val Lys Lys Ser 545 550 555 560

Ser Val Met Arg Glu Asn Glu Asn Gly Glu Glu Glu Glu Glu Glu Ala 565 570 575

Glu Phe Gly Glu Glu Asp Leu Phe His Gln Gln Gly Asp Pro Arg Thr 580 585 590

Thr Ser Arg Gly Cys Tyr Val Met 595 600

<210> 257

<211> 620

<212> PRT

<213> Homo sapien

<400> 257

Met Ser Pro Pro Ser Pro Gly Arg Arg Arg Glu Gln Arg Arg Pro Arg 1 5 10 15

- Ala Ala Ala Thr Met Ala Thr Pro Leu Pro Gly Arg Ala Gly Gly Pro 20 25 30
- Ala Thr Pro Leu Ser Pro Thr Arg Leu Ser Arg Leu Gln Glu Lys Glu 35 40 45
- Glu Leu Arg Glu Leu Asn Asp Arg Leu Ala His Tyr Ile Asp Arg Val 50 55 60
- Arg Ala Leu Glu Leu Glu Asn Asp Arg Leu Leu Leu Lys Ile Ser Glu 65 70 75 80
- Lys Glu Glu Val Thr Thr Arg Glu Val Ser Gly Ile Lys Ala Leu Tyr 85 90 95
- Glu Ser Glu Leu Ala Asp Ala Arg Arg Val Leu Asp Glu Thr Ala Arg 100 105 110
- Glu Arg Ala Arg Leu Gln Ile Glu Ile Gly Lys Leu Arg Ala Glu Leu 115 120 125
- Asp Glu Val Asn Lys Ser Ala Lys Lys Arg Glu Gly Glu Leu Thr Val 130 135 140
- Ala Gln Gly Arg Val Lys Asp Leu Glu Ser Leu Phe His Arg Ser Glu 145 150 155 160
- Val Glu Leu Ala Ala Ala Leu Ser Asp Lys Arg Gly Leu Glu Ser Asp 165 170 175
- Val Ala Glu Leu Arg Ala Gln Leu Ala Lys Ala Glu Asp Gly His Ala 180 185 190
- Val Ala Lys Lys Gln Leu Glu Lys Glu Thr Leu Met Arg Val Asp Leu 195 200 205
- Glu Asn Arg Cys Gln Ser Leu Gln Glu Glu Leu Asp Phe Arg Lys Ser 210 215 220
- Val Phe Glu Glu Glu Val Arg Glu Thr Arg Arg Arg His Glu Arg Arg 225 230 235 240
- Leu Val Glu Val Asp Ser Ser Arg Gln Gln Glu Tyr Asp Phe Lys Met 245 250 255
- Ala Gln Ala Leu Glu Glu Leu Arg Ser Gln His Asp Glu Gln Val Arg

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260 265 270 Leu Tyr Lys Leu Glu Leu Glu Gln Thr Tyr Gln Ala Lys Leu Asp Ser 280 Ala Lys Leu Ser Ser Asp Gln Asn Asp Lys Ala Ala Ser Ala Ala Arg 295 Glu Glu Leu Lys Glu Ala Arg Met Arg Leu Glu Ser Leu Ser Tyr Gln 310 315 Leu Ser Gly Leu Gln Lys Gln Ala Ser Ala Ala Glu Asp Arg Ile Arg 330 Glu Leu Glu Glu Ala Met Ala Gly Glu Arg Asp Lys Phe Arg Lys Met 340 345 Leu Asp Ala Lys Glu Gln Glu Met Thr Glu Met Arg Asp Val Met Gln 355 360 365 Gln Gln Leu Ala Glu Tyr Gln Glu Leu Leu Asp Val Lys Leu Ala Leu 375 Asp Met Glu Ile Asn Ala Tyr Arg Lys Leu Leu Glu Gly Glu Glu Glu 385 390 Arg Leu Lys Leu Ser Pro Ser Pro Ser Ser Arg Val Thr Val Ser Arg 405 410 Ala Thr Ser Ser Ser Ser Gly Ser Leu Ser Ala Thr Gly Arg Leu Gly 425 Arg Ser Lys Arg Lys Arg Leu Glu Val Glu Glu Pro Leu Gly Ser Gly Pro Ser Val Leu Gly Thr Gly Thr Gly Gly Ser Gly Gly Phe His Leu 455 Ala Gln Gln Ala Ser Ala Ser Gly Ser Val Ser Ile Glu Glu Ile Asp ` 470 475 Leu Glu Gly Lys Phe Val Gln Leu Lys Asn Asn Ser Asp Lys Asp Gln 490

Ser Leu Gly Asn Trp Arg Ile Lys Arg Gln Val Leu Glu Glu Glu 500 505 510

Ile Ala Tyr Lys Phe Thr Pro Lys Tyr Ile Leu Arg Ala Gly Gln Met 515 520 525

Val Thr Val Trp Ala Ala Gly Ala Gly Val Ala His Ser Pro Pro Ser 530 540

Thr Leu Val Trp Lys Gly Gln Ser Ser Trp Gly Thr Gly Glu Ser Phe 545 550 555 560

Arg Thr Val Leu Val Asn Ala Asp Gly Glu Glu Val Ala Met Arg Thr 565 570 575

Val Lys Lys Ser Ser Val Met Arg Glu Asn Glu Asn Gly Glu Glu Glu 580 585 590

Glu Glu Glu Ala Glu Phe Gly Glu Glu Asp Leu Phe His Gln Gln Gly 595 600 605

Asp Pro Arg Thr Thr Ser Arg Gly Cys Tyr Val Met 610 615 620

<210> 258

<211> 237

<212> PRT

<213> Homo sapien

<400> 258

Met Ser Pro Pro Ser Pro Gly Arg Arg Glu Gln Arg Arg Pro Arg

1 10 15

Ala Ala Ala Thr Met Ala Thr Pro Leu Pro Gly Arg Ala Gly Gly Pro
20 25 30

Ala Thr Pro Leu Ser Pro Thr Arg Leu Ser Arg Leu Gln Glu Lys Glu 35 40 45

Glu Leu Arg Glu Leu Asn Asp Arg Leu Ala His Tyr Ile Asp Arg Val 50 60

Arg Ala Leu Glu Leu Glu Asn Asp Arg Leu Leu Leu Lys Ile Ser Glu 65 70 75 80

Lys Glu Glu Val Thr Thr Arg Glu Val Ser Gly Ile Lys Ala Leu Tyr 85 90 95

Glu Ser Glu Leu Ala Asp Ala Arg Arg Val Leu Asp Glu Thr Ala Arg
100 105 110

Glu Arg Ala Arg Leu Gln Ile Glu Ile Gly Lys Leu Arg Ala Glu Leu 115 120 125

Asp Glu Val Asn Lys Ser Ala Lys Lys Arg Glu Gly Glu Leu Thr Val 130 135 140

Ala Gln Gly Arg Val Lys Asp Leu Glu Ser Leu Phe His Arg Ser Glu 145 150 155 160

Val Glu Leu Ala Ala Ala Leu Ser Asp Lys Arg Gly Leu Glu Ser Asp 165 170 175

Val Ala Glu Leu Arg Ala Gln Leu Ala Lys Ala Glu Asp Gly His Ala 180 185 190

Val Ala Lys Lys Gln Leu Glu Lys Gly Cys Pro Cys Ser Gln Lys Ser 195 200 205

Arg Ser His Val Asp Arg Gly Gly Arg Ile Leu Pro Lys His Phe Leu 210 215 220

Leu Glu Ala Thr Pro Leu Cys Ser Gln Ser Gly Gly Trp 225 230 235

<210> 259

<211> 620

<212> PRT

<213> Homo sapien

<400> 259

Met Ser Pro Pro Ser Pro Gly Arg Arg Glu Gln Arg Arg Pro Arg 1 5 10 15

Ala Ala Ala Thr Met Ala Thr Pro Leu Pro Gly Arg Ala Gly Gly Pro  $20 \hspace{1cm} 25 \hspace{1cm} 30$ 

Ala Thr Pro Leu Ser Pro Thr Arg Leu Ser Arg Leu Gln Glu Lys Glu 35 40 45

Glu Leu Arg Glu Leu Asn Asp Arg Leu Ala His Tyr Ile Asp Arg Val 50 55 60

Arg Ala Leu Glu Leu Glu Asn Asp Arg Leu Leu Leu Lys Ile Ser Glu 65 70 75 80

Lys Glu Glu Val Thr Thr Arg Glu Val Ser Gly Ile Lys Ala Leu Tyr 85 90 95

Glu Ser Glu Leu Ala Asp Ala Arg Arg Val Leu Asp Glu Thr Ala Arg 100 105 110

Glu Arg Ala Arg Leu Gln Ile Glu Ile Gly Lys Leu Arg Ala Glu Leu 115 120 125

Asp Glu Val Asn Lys Ser Ala Lys Lys Arg Glu Gly Glu Leu Thr Val 130 135 140

Ala Gln Gly Arg Val Lys Asp Leu Glu Ser Leu Phe His Arg Ser Glu 145 150 155 160

Val Glu Leu Ala Ala Ala Leu Ser Asp Lys Arg Gly Leu Glu Ser Asp 165 170 175

Val Ala Glu Leu Arg Ala Gln Leu Ala Lys Ala Glu Asp Gly His Ala 180 185 190

Val Ala Lys Lys Gln Leu Glu Lys Glu Thr Leu Met Arg Val Asp Leu 195 200 205

Glu Asn Arg Cys Gln Ser Leu Gln Glu Glu Leu Asp Phe Arg Lys Ser 210 220

Val Phe Glu Glu Glu Val Arg Glu Thr Arg Arg Arg His Glu Arg Arg 225 230 235 240

Leu Val Glu Val Asp Ser Ser Arg Gln Gln Glu Tyr Asp Phe Lys Met 245 250 255

Ala Gln Ala Leu Glu Glu Leu Arg Ser Gln His Asp Glu Gln Val Arg 260 265 270

Leu Tyr Lys Leu Glu Leu Glu Gln Thr Tyr Gln Ala Lys Leu Asp Ser 275 280 285

Ala Lys Leu Ser Ser Asp Gln Asn Asp Lys Ala Ala Ser Ala Ala Arg 290 295 300

Glu Glu Leu Lys Glu Ala Arg Met Arg Leu Glu Ser Leu Ser Tyr Gln 305 310 315 320 WO 2004/050900 PCT/US2003/040131

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Leu Ser Gly Leu Gln Lys Gln Ala Ser Ala Ala Glu Asp Arg Ile Arg 325 Glu Leu Glu Glu Ala Met Ala Gly Glu Arg Asp Lys Phe Arg Lys Met Leu Asp Ala Lys Glu Gln Glu Met Thr Glu Met Arg Asp Val Met Gln 360 Gln Gln Leu Ala Glu Tyr Gln Glu Leu Leu Asp Val Lys Leu Ala Leu 370 375 Asp Met Glu Ile Asn Ala Tyr Arg Lys Leu Leu Glu Gly Glu Glu Glu 390 395 Arg Leu Lys Leu Ser Pro Ser Pro Ser Ser Arg Val Thr Val Ser Arg 405 Ala Thr Ser Ser Ser Ser Gly Ser Leu Ser Ala Thr Gly Arg Leu Gly 425 Arg Ser Lys Arg Lys Arg Leu Glu Val Glu Glu Pro Leu Gly Ser Gly 440 Pro Ser Val Leu Gly Thr Gly Thr Gly Gly Ser Gly Gly Phe His Leu 455 Ala Gln Gln Ala Ser Ala Ser Gly Ser Val Ser Ile Glu Glu Ile Asp Leu Glu Gly Lys Phe Val Gln Leu Lys Asn Asn Ser Asp Lys Asp Gln 490 Ser Leu Gly Asn Trp Arg Ile Lys Arg Gln Val Leu Glu Gly Glu Glu 500 505 Ile Ala Tyr Lys Phe Thr Pro Lys Tyr Ile Leu Arg Ala Gly Gln Met 520 515 Val Thr Val Trp Ala Ala Gly Ala Gly Val Ala His Ser Pro Pro Ser 540

Thr Leu Val Trp Lys Gly Gln Ser Ser Trp Gly Thr Gly Glu Ser Phe

550

555 560

Arg Thr Val Leu Val Asn Ala Asp Gly Glu Glu Val Ala Met Arg Thr 565 570 . 575

Val Lys Lys Ser Ser Val Met Arg Glu Asn Glu Asn Gly Glu Glu Glu 580 585 590

Glu Glu Glu Ala Glu Phe Gly Glu Glu Asp Leu Phe His Gln Gln Gly 595 600 605

Asp Pro Arg Thr Thr Ser Arg Gly Cys Tyr Val Met 610 615 620